

FIG. I

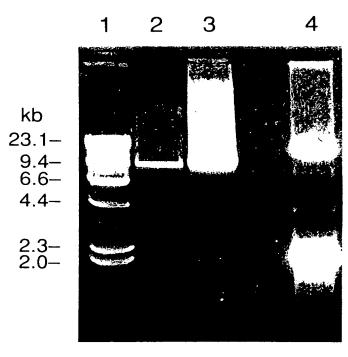


FIG. 2

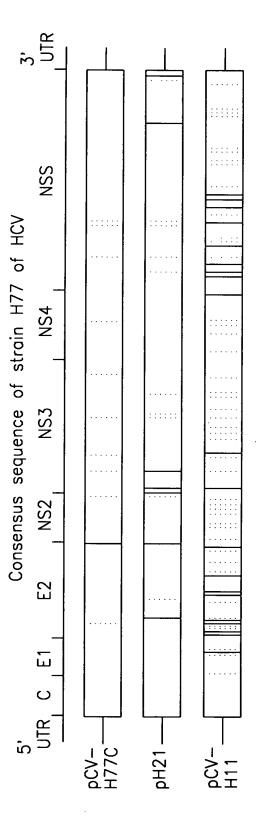


FIG. 3

				
10 20	30	40	50	
<u>1234567890 1234567890</u>	<u>1234567890</u>	1234567890	1234567890	•
GCCAGCCCCC TGATGGGGC	GACACTCCAC	CATGAATCAC	TCCCCTGTGA	50
GGAACIACIG TCITCACGCA	GAAAGOGICT	AGCCATGGCG	TTAGTATGAG	100
TGICGIGCAG CCTCCAGGAC	CCCCCCICCC	GGGAGAGCCA	TAGICGICIG	150
CGGAACCGGT GAGTACACCG	GAATTGCCAG	GACGACCGGG	TCCTTTCTTG	200
GATAAACCCG CŢCAATGCCT	GGAGATTTGG	COCLIGACOCC	GCAAGACTGC	250
TAGCCGAGIA GIGITGGGIC	GCGAAAGGCC	TIGIGGIACT	GCCTGATAGG	300
GIGCTIGCGA GIGCCCCGGG	AGGICICGIA	GACCGIGCAC	CATGAGCACG	350
AATCCIAAAC CTCAAAGAAA	AACCAAACGT	AACACCAACC	GICGCCCACA	400
GGACGICAAG TICCCGGGIG	GOGGICAGAT	CGITGGIGGA	GITTACTIGT	450
TGCCGCGCAG GGGCCCTAGA	TIGGGIGIGC	GCGCCACGAG	GAAGACTTCC	500
GAGCGGTCGC AACCTCGAGG	TAGACGICAG	CCTATCCCCA	AGGCACGTCG	550
GCCCGAGGGC AGGACCTGGG	CTCAGCCCGG	GIACCCITIGG	CCCCICIAIG	600
GCAATGAGGG TTGCGGGTGG	GCGGGATGGC	TCCIGICICC	CCCIGCCICT	650
CGGCCTAGCT GGGGCCCCAC	AGACCCCCGG	CGTAGGTCGC	GCAATTIGGG	700
TAACGICATC GATACCCTTA	CGIGCGGCIT	CCCCGACCIC	ATGGGGTACA	750
TACCOCTCGT COCCOCCCT	CTTGGAGGCG	CTGCCAGGGC	CCTGGCGCAT	800
GGCGICCGGG TICIGGAAGA	CGGCGIGAAC	TATGCAACAG	GCAACCTTCC	850
TEGITECICI TICICIATET	TCCTTCTGGC	CCIGCICICI	TOCCIGACIG	900
TGCCCGCTTC AGCCTACCAA	GIGCGCAATT	CCICGGGGCT	TTACCATGIC	950
ACCAATGATT GCCCTAACTC	GAGIATIGIG	TACGAGGCGG	CCGATGCCAT	1000
CCTGCACACT CCGGGGTGTG	TCCCTTGCGT	TCGCGAGGGT	AACGCCTCGA	1050
GGIGIIGGGI GGCGGIGACC	CCCACGGIGG	CCACCAGGGA	CGGCAAACTC	1100
CCCACAACGC AGCTTCGACG	TCATATCGAT	CICCIIGICG	GGAGCGCCAC	1150
CCTCTGCTCG GCCCTCTACG	TGGGGGACCT	GIGCGGGICI	GICTITCITG	1200
TIGGICAACT GITTACCTIC	TCTCCCAGGC	GCCACTGGAC	GACGCAAGAC	1250
TGCAATIGIT CIATCIATCO	CGGCCATATA	ACGGGICATC	GCATGGCATG	1300
OGATATGATG ATGAACTOGT	, CCCCLYCCCC	ACCETTICETE	GIAGCICAGC	1350
TGCTCCGGAT CCCACAAGCC	ATCATGGACA	TGATCGCTGG	TGCTCACTGG	1400
GGAGTCCTGG CGGGCATAGC	GIATTICICC	ATGGTGGGGA	ACTGGGCGAA	1450
GGICCIGGIA GIGCIGCIG	TATTIGCOGG	CCTCCACCCC	GAAACCCACG	1500
TCACCGGGG AAATGCCGGC	CGCACCACGG	CIGGGCTIGI	TEGICICCIT	1550
ACACCAGGCG CCAAGCAGAA	CATCCAACIG	ATCAACACCA	ACGGCAGTTG	1600
GCACATCAAT AGCACGGCCT	TGAATTGCAA	TGAAAGCCIT	AACACCGGCT	1650
GGTTAGCAGG GCTCTTCTAT	CAACACAAAI	TCAACICITC	AGGCIGICCT	1700
CACACCTICC CCACCTICCC				1750
TCCTATCAGT TATGCCAACC	GAAGCGGCCI	CGACGAACGC	CCCIACIGCT	1800
GCACTACCC TCCAAGACC	TGIGGCATIC	TGCCCGCAAA	GAGCGIGIGI	1850
GGCCCGGIAT ATTGCTTCAL	C TOOCAGOOO	् दाव्हाव्हाव्ह	GAACGACCGA	1900

FIG. 4A

10	20	30	40	50	
			1234567890		
			TGCAAATGAT		1950
			GCAATIGGIT		2000
			TGCGGAGCGC		2050
CATCGGAGGG			CIGCCCCACT		2100
GCAAACATCC			GOGGCICCGG		2150
ACACCCAGGI	GCATGGTCGA	CTACCCGUAT	AGGCTTTGGC	ACTATCCTTG	2200
			GATGIACGIG		2250
AGCACAGGCT	GGAAGOGGCC	TGCAACTGGA	CGCCGCGCCA	ACCCIGIGAT	2300
CIGGAAGACA	GGGACAGGIC	CGAGCICAGC	COGTICCICC	TGTCCACCAC	2350
ACAGIGGCAG	GICCLICCGL	GITCITICAC	GACCCTGCCA	GCCTTGTCCA	2400
CCCCCCTCAT	CCACCTCCAC	CAGAACATIG	TGGACGTGCA	GEACTIGEAC	2450
GCGCTACCGT	CAAGCATCGC	GICCIGGGCC	ATTAAGTGGG	AGTACGTCGT	2500
TCTCCTGTTC	CTTCTCCTTG	CAGACGCGCG	CGICICCICC	TGCTTGTGGA	2550
TGATGITACT	CATATCCCAA	CCCCACCCCC	CTTTCGAGAA	CCTCGTAATA	2600
CICAAIGCAG	CATCCCTGGC	CGGGACGCAC	GCCTTGTGT	CCTTCCTCGT	2650
GITCITCICC	TTTGCGTGGT	ATCTGAAGGG	TAGGIGGGIG	CCCGGAGGGG	2700
TCTACGCCCT	CTACGGGATG	TEGECTETEC	TCCTGCTCCT	GCTGGCGTTG	2750
CCTCAGCGGG	CATACGCACT	GGACACGGAG	GIGGCCGCGT	CGIGIGGCCG	2800
CGTTGTTCTT	GICGGGTTAA	TGGCGCTGAC	TCTGTCGCCA	TATTACAAGC	2850
GCTATATCAG	CTGGTGCATG	TEGIESCITC	AGTATTTTCT	GACCAGAGTA	2900
GAAGCGCAAC	TCCACGIGIG	GGTTCCCCCCC	CTCAACGTCC	GGGGGGGGG	2950
CGATCCCGTC	ATCTTACTCA	TGIGIGIAGT	ACACCCGACC	CIGGIATTIG	3000
ACATCACCAA	ACIACICCIG	GCCATCITCG	CACCCCTTTG	CATTCTTCAA	3050
GCCAGTTTGC	TTAAAGICCC	CIACTICGIG	CGCGTTCAAG	CCCTTCTCCCG	3100
GATCTGCGCG	CTAGCGCGGA	AGATAGCCGG	AGGICATTAC	GTGCAAATGG	3150
CCATCATCAA	GITIAGGGGGG	CITACIGGCA	CCIAIGIGIA	TAACCATCIC	3200
ACCCCICITC	GAGACTGGGC	GCACAACGGC	CIGOGAGAIC	TEGCCGTEGC	3250
TGTGGAACCA	GIOGICTICT	CCCCAATGCA	GACCAAGCTC	ATCACGTGGG	3300
GGGCAGATAC	CCCCCCCTTCC	GGTGACATCA	TCAACGGCTT	GCCCGTCTCT	3350
CCCCGTAGGG	CCCACCACAT	ACTOCTTOGG	CCACCCGACG	CAATOGICIC	3400
CAAGGGGTGG	AGGITGCIGG	CGCCCATCAC	GGCGTACGCC	CAGCAGACGA	3450
CACCCTCCT	AGGGIGIATA	ATCACCAGCC	TGACTGGCCG	GCACAAAAAC	3500
CAAGTGGAGG	GIGAGGICCA	GATCGIGICA	ACTGCTACCC	AAACCTTCCT	3550
GGCAACGTGC	ATCAATGGGG	TATGCTGGAC	TGTCTACCAC	GGGGCCGGAA	3600
CGAGGACCAT	CGCATCACCC	AAGGGTCCTG	TCATCCAGAT	GIATACCAAT	3650
GTGGACCAAG	ACCTTGTGGG	CIGGCCCCCT	CCTCAAGGIT	CCCGCTCATT	3700
GACACCCTGT	ACCIGCGGCT	CCTCGGACCT	TIACCIGGIC	ACCACCCACC	3750
CCGATGTCAT	TCCCGTCCCC	CCCCCACCTG	ATAGCAGGGG	TAGCCIGCIT	3800

FIG. 4B

					
10	20	30	40	50	
	1234567890				
	CCATTICCIA				3850
	GGACACGCCG			· · ·	3900
	TAAAGCGGTG			-	3950
	CCCCGGIGIT				4000
	CAGGIGGCCC				4050
	CCCGGCIGCG				4100
	CIGITICCICC				4150
	GITGATCCIA				4200
	CATCACGIAC	•			4250
	GAGGIGCITA				4300
CACGGATGCC	ACATCCATCT	TGGGCATCGG	CACIGICCIT	GACCAAGCAG	4350
AGACTGCGGG	GGCGAGACTG	GLIGIGGIGG	CCACIGCIAC	CCCICCGGGC	4400
TCCGTCACTG	TGTCCCATCC	TAACATOGAG	GAGGITICCIC	TGTCCACCAC	4450
CCCACACATC	CCCTTTIACG	GCAAGGCTAT	CCCCCTCCAG	GIGATCAACG	4500
GGGGAAGACA	TCTCATCTTC	TGCCACTCAA	AGAAGAAGIG	CCACCACCTC	4550
GCCGCGAAGC	TOGTOGCATT	GGGCATCAAT	CCCGTCCCCT	ACTACCGCGG	4600
TCTTGACGIG	TCTGTCATCC	CGACCAGCGG	CGATGITGIC	GICGIGICGA	4650
CCGATGCTCT	CATGACTGGC	TTTACCGGCG	ACTICGACIC	TGTGATAGAC	4700
TGCAACACGT	GIGICACICA	GACAGTCGAT	TICAGCCIIG	ACCCTACCTT	4750
TACCATIGAG	ACAACCACGC	TCCCCCAGGA	TECTETCTCC	AGGACTCAAC	4800
GCCGGGGCAG	GACTGGCAGG	GGGAAGCCAG	GCATCTATAG	ATTIGIGGCA	4850
CCGGGGGAGC	GCCCTCCGG	CATGITCGAC	TOGICOGICC	TCTGTGAGTG	4900
CTATGACGCG	GCTGTGCTT	GGTATGAGCT	CACGCCCCCC	GAGACTACAG	4950
TTAGGCTACG	ACCGIACATG	AACACCCCGG	GCTTCCCCT	GTGCCAGGAC	5000
CATCTTGAAT	TITTEGGAGGG	CGICTTIACG	GGCCTCACTC	ATATAGATGC	5050
CCACTTTTA	TCCCAGACAA	ACCAGAGICG	GCAGAACTTT	CCTTACCTGG	5100
TAGCGIACCA	AGCCACCGIG	TGCGCTAGGG	CICAAGCCCC	TCCCCCATCG	5150
TGGGACCAGA	TGTGGAAGTG	TTTGATCCCC	CITIAAACCCA	CCCTCCATGG	5200
GCCAACACCC	CIGCIATACA	GACTGGGGGC	TGTTCAGAAT	GAAGTCACCC	5250
TGACGCACCC	AATCACCAAA	TACATCATGA	CATGCATGIC	GGCCGACCTG	5300
GAGGICGICA	CGAGCACCTG	GGIGCICGIT	GEOGGETICC	TESCIGCICT	5350
GGCCGCGTAI	TGCCTGTCAA	CAGGCTGCGT	GCICATAGIG	GCCAGGATCG	5400
TCTTGTCCCC	GAAGCCGGCA	ATTATACCIG	ACAGGGAGGT	TCTCTACCAG	5450
GAGTTCGATC	AGATOGAAGA	GIGCICICAG	CACTTACCGT	ACATOGAGCA	5500
ACCGATGATO	CICCCICACC	AGITCAAGCA	GAAGGCCCTC	GECCICCIEC	5550
AGACCGCGTC	CCGCCATGCA	GAGGITATCA	CCCTCCTGT	CCAGACCAAC	5600
TOGCAGAAAC	CTCGAGGICIT	TICCCCAAC	CACATGTGGA	ATTTCATCAG	5650
TOGGATACA	A TACTTOGCOC	GCCTGTCAAC	CIGCCIGGI	' AACCCCGCCA	5700

FIG. 4C

					
10	20	30	40	50	
<u>1234567890</u>			1234567890		
	GATGGCTTTT				5750
GGCCAAACCC					5800
accaccacac			GGGIGCIGGC		5850
CCGCCATCGG	CAGCGITIGGA	CTGGGGAAGG	TCCTCGTGGA	CATTCTTGCA	5900
GGGTATGGCG	CEGGCGIGGC	GGGAGCTCTT	GIAGCATTCA	AGATCATGAG	5950
CGGTGAGGTC	CCCTCCACCG	AGGACCIGGI	CAATCIGCIG	CCCCCATCC	6000
TCTCGCCTGG	AGCCCTIGIA	GICGGIGIGG	TCIGCGCAGC	AATACTGCGC	6050
CGGCACGITG	GCCCGGGGGGA	GGGGGCAGTG	CAATGGATGA	ACCGCTAAT	6100
AGCCTTCGCC	TCCCGGGGGGA	ACCATGITIC	CCCCACGCAC	TACGIGCCGG	6150
AGAGCGATGC	AGCCGCCCCCC	GICACIGOCA	TACTCAGCAG	CCTCACTGIA	6200
ACCCAGCICC	TGAGGCGACT	CCATCAGICG	ATAAGCTCGG	AGIGIACCAC	6250
TCCATGCTCC	GGTTCCTGGC	TAAGGGACAT	CTGGGACTGG	ATATGCGAGG	6300
TGCTGAGCGA	CITTAAGACC	TGGCTGAAAG	CCAAGCTCAT	GCCACAACTG	6350
CCTGGGATTC	CCTTTGTGTC	CTGCCAGCGC	GGGTATAGGG	GGGTCTGGCG	6400
AGGAGACGGC	ATTATGCACA	CTCGCTGCCA	CIGIGGAGCT	GAGATCACTG	6900
GACATGTCAA	AAACGGGACG	ATGAGGATCG	TCGGTCCTAG	GACCTGCAGG	6950
AACATGTGGA	GIGGGACGIT	CCCCATTAAC	GCCTACACCA	CCCCCCTG	6550
TACTCCCCTT	CCTGCGCCGA	ACTATAAGTT	CCCCTCTCC	AGGGIGICIG	6600
CAGAGGAATA	CGIGGAGATA	AGGCGGGTGG	GGGACTTCCA	CTACGIATCG	6650
GGTATGACTA	CIGACAAICT	TAAATGCCCG	TGCCAGATCC	CATCGCCCGA	6700
ATTTTTCACA	GAATTGGACG	GGGTGCGCCT	ACACAGGITT	GCGCCCCTT	6750
GCAAGCCCTT	GCTGCGGGAG	GAGGIAICAT	TCAGAGTAGG	ACTOCACGAG	6800
TACCCCGTCG	CGTCCCAATT	ACCTTGCGAG	CCCGAACCGG	ACGIAGCCGI	6850
GIIGACGICC	ATGCTCACTG	ATCCCTCCCA	TATAACAGCA	CAGGGGGGCCG	6900
GGAGAAGGTT	CCCCACACACCC	TCACCCCTT	CTATGGCCAG	CICCICGGCT	6950
AGCCAGCIGI	CCCCTCCATC	TCTCAAGGCA	ACTTGCACCG	CCAACCATGA	7000
CICCCCIGAC	GCCGAGCTCA	TAGAGGCTAA	CCTCCTGTGG	AGGCAGGAGA	7050
TGGGCGGCAA	CATCACCAGG	GTTGAGTCAG	AGAACAAAGT	GGIGATICIG	7100
GACTCCTTCG	ATCCGCTTGT	GGCAGAGGAG	CATCACCCCC	AGGICICCGT	7150
ACCTGCAGAA	ATTCTGCGGA	AGTCTCGGAG	ATTCGCCCGG	CCCTGCCCG	7200
TCTGGGGGG	GCCGGACTAC	AACCCCCCCCC	TAGTAGAGAC	GIGGAAAAAG	7250
CCTGACTACG	AACCACCIGI	GGICCATGGC	TGCCCGCTAC	CACCTCCACG	7300
GICCCCICCI	GIGCCICCGC	CTCGGAAAAA	GOGTACOGTG	GICCICACCG	7350
AATCAACCCT	ATCTACTGCC	TTGGCCGAGC	TTGCCACCAA	AAGITTTOGC	7400
AGCTCCTCAA	CTTCCGGCAT	TACGGGGGAC	AATACGACAA	CATCCTCTGA	7450
CCCCCCCCCT	TCTGGCTGCC	CCCCCGACTC	CCACCITICAG	TCCTATTCTT	7500
CCATGCCCCC	CCTGGAGGGG	GAGCCTGGGG	ATCCCGATCT	CAGCGACGGG	7550
TCATCGTCGA	COGTCAGTAG	TGGGGCCGAC	ACCGAAGATG	TOGTGTGCTG	7600

			· 		
10	20	30	40	50	
1234567890	<u>1234567890</u>	<u>1234567890</u>	1234567890	1234567890	··
CICAAIGICI	TATICCIGGA	CAGGCGCACT	CGTCACCCCG	TECCCTCCCC	7650
AAGAACAAAA	ACTGCCCATC	AACGCACTGA	GCAACICGIT	GCTACGCCAT	7700
CACAATCIGG	TGTATTCCAC	CACTICACGC	AGIGCTIGCC	AAAGGCAGAA	7750
GAAAGICACA	TTTGACAGAC	TGCAAGITCT	GGACAGCCAT	TACCAGGACG	7800
TGCTCAAGGA	GGTCAAAGCA	GCGGCGTCAA	AAGIGAAGGC	TAACITGCIA	7850
TCCGTAGAGG	AAGCTTGCAG	CCTGACGCCC	CCACATTCAG	CCAAATCCAA	7900
GITTGGCTAT	GGGGCAAAAG	ACGICCGITG	CCATCCCAGA	AAGGCCGTAG	7950
CCCACATCAA	CICCGIGICG	AAAGACCITC	TGGAAGACAG	TGTAACACCA	8000
ATAGACACTA	CCATCATGGC	CAAGAACGAG	GITTICIGCG	TICAGCCIGA	8050
CAACCCCCT	CGTAAGCCAG	CICGICICAT	CCICITOCCC	CACCICCCC	8100
TECECGIGIG	CGAGAAGAIG	GCCCTGTACG	ACGIGGITAG	CAAGCTCCCC	8150
CTGGCCGTGA	TGGGAAGCIC	CTACGGATTC	CAATACTCAC	CAGGACAGCG	8200
GGTTGAATTC	CTCGTGCAAG	CGTGGAAGTC	CAAGAAGACC	CCGATGGGGT	8250
TCTCGTATGA	TACCCCCTGT	TTTGACTCCA	CAGICACIGA	GAGCGACATC	8300
CGTACGGAGG	AGGCAATTTA	CCAATGITGI	GACCIGGACC	CCCAAGCCCG	8350
CGTGGCCATC	AAGTCCCTCA	CTGAGAGGCT	TTATGITGGG	GGCCCTCTTA	8400
CCAATTCAAG	GGGGGAAAAC	TGCGGCTACC	GCAGGTGCCG	CCCCACCCCC	8450
GTACTGACAA	CTACCTCTCC	TAACACCCTC	ACTIGCTACA	TCAAGGCCCG	8500
GGCAGCCTGT	CGAGCCGCAG	GGCTCCAGGA	CTGCACCATG	CICCICICIC	8550
GCGACGACTT	' AGICGITAIC	TGIGAAAGIG	CGGGGGTCCA	GGAGGACGCG	8600
GCGAGCCTGA	GAGCCTTCAC	GGAGGCTATG	ACCAGGIACT	000000000000000000000000000000000000000	8650
CGGGGACCCC	CCACAACCAG	AATACGACTT	GGAGCTTATA	ACATCATGCT	8700
CCTCCAACGI	GICAGICGCC	CACGACGGGG	CTGGAAAGAG	GGICIACIAC	8750
CTTACCCGTG	ACCCTACAAC	CCCCCTCCCCC	AGAGCCGCGT	GGGAGACAGC	8800
AAGACACACT	CCAGICAATI	' CCTGGCTAGG	CAACATAATC	ATGTTTGCCC	8850
CCACACTGTG	GCCACCATC	ATACIGATGA	CCCATTICIT	TAGCGICCIC	8900
ATAGCCAGGC	ATCAGCTIGA	ACAGGCTCTT	AACTGTGAGA	TCTACGGAGC	8950
CIGCIACICO	ATAGAACCAC	TOGATCTACC	TCCAATCATT	CAAAGACTCC	9000
ATGGCCTCAC	COCATTITCA	CICCACAGII	· ACTCTCCAGG	TGAAATCAAT	9050
AGGTGGCCC	CATGCCTCAC	AAAACTTGGG	GICCCCCCCI	TOCCACCTTC	9100
GAGACACCGC	GCCCGGAGCC	TOCGOCTAC		AGAGGAGGCA	9150
GGGCTGCCAT	r atgrogcaac	TACCICITO	ACTOGGCAGI	' AAGAACAAAG	9200
CICAAACIC	A CTCCAATAG	C GGCCGCTGGC	COGCTOGACT	TGICCOGITIG	9250
GITCACGGC.	r ggctacagco	G GGGGAGACAI	TTATCACAGO	GIGICICAIG	9300
ccccccccc	G CIGGITCIG	TITIOCCIAC	C TCCTGCTCGC	TGCAGGGGTA	9350
GGCATCTAC	C TOCTOCCAN	A CCGATGAAG	G TIGGGGIAAA	A CACTCCGGCC	9400
TCTTAAGCC	A TITOCIGIT	r trittritti	r trrttrttt	TITTICITIT	9450
TTTTTTCT	r tectricer	r crrrrrra	TTTCTTTTTC	CCTTCTTTAA	9500

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TEGTESCTCC	ATCTTAGCCC	TAGTCACGGC	TAGCIGIGAA	AGGICCGIGA	9550
GCCGCATGAC	TCCAGAGAGT	GCTGATACTG	GCCTCTCTGC	AGATCATGT	9599

FIG. 4F

10	20	30	40	50	
<u>1234567890</u>	1234567890	1234567890	1234567890	1234567890	·
MSINPKPQRK	TKRNINRRPQ	DVKFPGGGQI	VGGVYLLPRR	GPRLGVRATR	50
~	RRQPIPKARR	-			100
RGSRPSWGPT	DPRRRSRNLG	KVIDILICGF	ADLMGYIPLV	GAPLOGAARA	150
LAHGVRVLED	GVNYATGNLP	GCSFSIFLLA	LLSCLTVPAS	AYQVRNSSGL	200
YHVINDCPNS	SIVYEAADAI	LHIPGCVPCV	REGNASROW	AVIPIVATRD	250
CKLPITQLRR	HIDLLVGSAT	LCSALYVGDL	CGSVFLVGQL	FIFSPRRHWT	300
TODONOSIYP	CHITCHRMAW	DMMNWSPIA	ALVVAQLLRI	POATMOMIAG	350
AHWGVLAGIA	YFSMVGNWAK	VLVVLLLFAG	VDAETHVIGG	NAGRITAGLV	400
GLLTPGAKQN	IQLININGSW	HINSTALNON	ESLNIGWLAG	LFYOHKFNSS	4 50
GCPERLASCR	RLTDFAQGWG	PISYANGSGL	DERPYCWHYP	PRPCGIVPAK	500
SVCGPVYCFT	PSPVVVGITID	RSGAPTYSWG	ANDIDVFVLN	NIRPPLGNWF	550
GCTWMNSTGF	TKVCGAPPCV	IGGVGNNTLL	CPIDCFRKHP	EATYSROGSG	600
PWITPROMVD	YPYRLWHYPC	TINYTIFKVR	MYVGGVEHRL	EAACIWIRGE	650
RCDLEDRDRS	ELSPLLLSTT	QWQVLPCSFT	TLPALSTGLI	HLHQNIVDVQ	700
YLYGVGSSIA	SWAIKWEYVV	LLFLLLADAR	VCSCLWMMLL	ISQAEAALEN	750
LVILNAASLA	GIHGLVSFLV	FFCFAWYLKG	RWPGAVYAL	YGMWPLLLLL	800
LALPQRAYAL	DTEVAASCGG	VVLVGLMALT	LSPYYKRYIS	WOMWLQYFL	850
TRVEAQLHVW	VPPLNVRGGR	DAVILLMCVV	HPILVFDIIK	LLLAIFGPLW	900
ILQASLLKVP	YFVRVQGLLR	ICALARKIAG	CHYVQMAIIK	LGALTGTYVY	950
NHLTPLRDWA	HNGLRDLAVA	VEPVVFSRME	TKLITWGADI	AACCEDIINGL	1000
PVSARRQQEI	LLGPADGMVS	KGWRLLAPIT	AYAQQIRGLL	CCITTSLICER	1050
DKNQVEGEVQ	IVSTATQIFL	ATCINGVCWT	VYHGAGIRII	ASPKGPVIQM	1100
YTNVDQDLVG	WPAPQGSRSL	TPCTCGSSDL	YLVTRHADVI	PVRRRGDSRG	1150
SLLSPRPISY	LKGSSGGPLL	CPAGHAVGLF	RAAVCIRGVA	KAVDFIPVEN	1200
LGTIMRSPVF	TDNSSPPAVP	QSFQVAHLHA	PIGSGKSTKV	PAAYAAQGYK	1250
VLVLNPSVAA	TLGFGAYMSK	AHGVDPNIRT	GVRTTTTGSP	ITYSTYCKFL	1300
ADGGCSGGAY	DILICOECHS	TDATSILGIG	TVLDQAETAG	ARLVVLATAT	1350
PPGSVIVSHP	NIEEVALSTT	GEIPFYGKAI	PLEVIKGGRH	LIFCHSKKKC	1400
DELAAKLVAL	GINAVAYYRG	LDVSVIPISG	DVVVVSIDAL	MIGFIGDFDS	1450
VIDONICVIO	TVDFSLDPTF	TIETTTLPQD	AVSRTQRRGR	TGRGKPGIYR	1500
		· -		AYMNTPGLPV	1550
CQDHLEFWEC	VFIGLTHIDA	HFLSQTKQSG	ENFPYLVAYQ	ATVCARAQAP	1600
PPSWDQMWKC	LIRLKPTLHG	PIPLLYRLGA	VONEVILIHP	ITKYIMICMS	1650
ADLEVVISIV	VLVGGVLAAL	AAYCLSIGCV	VIVŒRIVLSG	KPAIIPDREV	1700
LYQEFDEMEE	CSQHLPYIEQ	GMMLAEQFKQ	KALGLIQTAS	RHAEVITPAV	1750
QIIVWQKLEVE	WAKHMWNFIS	GIQYLAGLSI	LPGNPALASL	MAFTAAVISP	1800
LTTGQTLLFN	ILGGWVAAQL	AAPGAATAFV	GAGLAGAAIG	SVGLGKVLVD	1850
ILAGYGAGVA	A GALVAFKIMS	GEVPSTEDLV	NLLPATLSPO	ALVVGVVCAA	1900

10	20	30	40	50	
	1234567890				
	GAVQWMNRLI				1950
LIVIQLLRRL	HOWISSECTT	PCSGSWLRDI	WDWICEVLSD	FKIWLKAKLM	2000
PQLPGIPFVS	CQRGYRGVWR	GDGIMHIRCH	CGAEITGHVK	NGIMRIVGPR	2050
TCRNMWSGIF	PINAYTIGPC	TPLPAPNYKF	ALWRVSAEEY	VEIRRVCDFH	2100
YVSGMITIDNL	KCPCQIPSPE	FFTELDGVRL	HRFAPPCKPL	LREEVSFRVG	2150
LHEYPVGSQL	PCEPEPDVAV	LTSMLTDPSH	ITAEAAGRRL	ARGSPPSMAS	2200
SSASQLSAPS	LKATCTANHD	SPDAELIEAN	LLWRQEMGGN	TTRVESENKV	2250
VILDSFDPLV	AEEDEREVSV	PAETLRKSRR	FARALPWAR	PDYNPPLVET	2300
WKKPDYEPPV	VHGCPLPPPR	SPPVPPPRKK	RIVVLIESIL	STALAFLATK	2350
SFGSSSTSGI	TGENTTTSSE	PAPSGCPPDS	DVESYSSMPP	LEGEPGDPDL	2400
SDGSWSTVSS	GADIEDVVCC	SMSYSWIGAL	VIPCAAEEQK	LPINALSNSL	2450
LRHHNLVYST	TSRSACQRQK	KVIFDRLQVL	DSHYQDVLKE	VKAAASKVKA	2500
NLLSVEEACS	LTPPHSAKSK	FGYGAKDVRC	HARKAVAHIN	SWKDLLEDS	2550
VIPIDITIMA	KNEVFCVQPE	KGGRKPARLI	VFPDLGVRVC	EKMALYDVVS	2600
KLPLAVMGSS	YGFQYSPGQR	VEFLVQAWKS	KKTPMGFSYD	TRCFDSIVIE	2650
SDIRTEEAIY	QCCDLDPQAR	VAIKSLTERL	YVGGPLINSR	GENCGYRRCR	2700
ASGVLTTSCG	NTLICYIKAR	AACRAAGLQD	CIMLVCGDDL	VVICESAGVQ	2750
	EAMIRYSAPP				2800
VYYLTRDPTT	PLARAAWETA	RHTPVNSWLG	NIIMFAPILW	ARMILMIHEF	2850
SVLIARDQLE	QALNCEIYGA	CYSIEPLDLP	PIIQRLHGLS	AFSLHSYSPG	2900
	KLGVPPLRAW		-		2950
RIKLKLTPIA	AAGRLDLSGW	FTAGYSGGDI	YHSVSHARPR	WFWFCLLLLA	3000
AGVGIYLLPN	R				3011

FIG. 4H

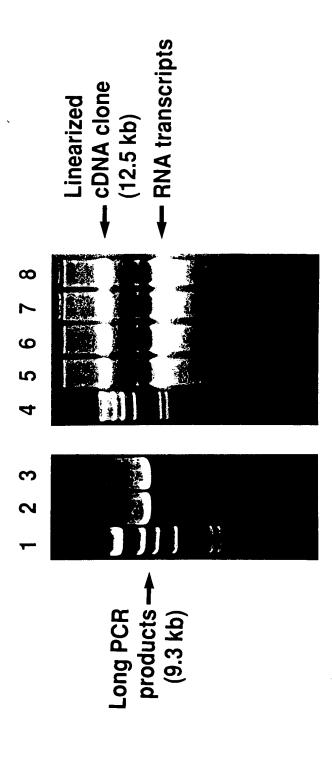


FIG. 5

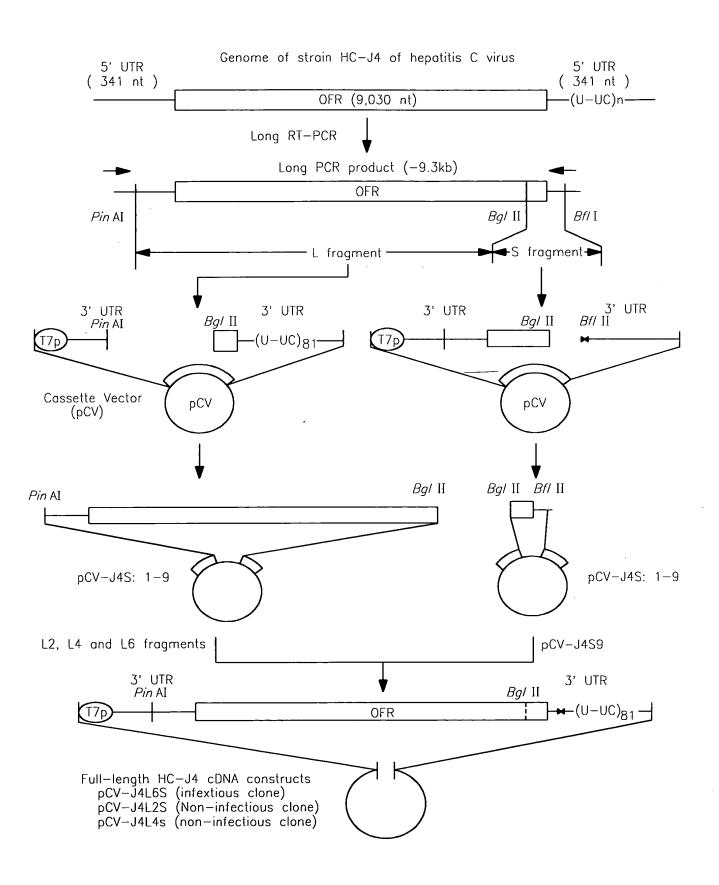


FIG. 6

_	_	·	т	т_	F	,	_		1				
Cons-F	z		ΑT	R.O	A	2	· 02	9	2			ı co	A
Cons-D			<u></u>	R.0			•	•			•		
(C)				•		•							
L10(B)		•	 	O		T	•	A					 -
(B) [1			<u> </u>	O	<u>-</u>	•		A	۵		•		•
L3(B)			_	O		I		A	۵				-
(A)67											⋖		
L8(A)		۵					o				•		
(A)91				•	•	•	o						
L2(A)	•			•	•	•	Ŏ		•				
L1* (A)	S			•	•	•	•		•	•	A		
Cons-p9	Z	7	А	R	А	æ	R	9	Z	Z	ш	ပ	¥
L fragment	16	36	52	70	189	195	231	233	234	250	567	304	379
	Core					E							

FIG. 7A

											_			_				_	,				
Cons-F	A	Y,H	S'1	R,G	۸	A,V	工	S	H,O	F,L	A,T	S	S	A,V	Y	K,E	I'A	^	۸			Ö	A
Cons-D	F,T	Y,H	T,S	. 9	•	^	•	•	エ		1	•	•	A,V	•]	I	•		•	•	•	•
(C)	•	•		•		^	2	•				Ь	•		•	E]]	•	•	•	•	•	•
L10(B)	L	٨	S	9	A	•	R		H		⊥	•	٠	٨	٠	Ŀ	1	•	•	•	•	•	•
(B) _* (7)	1	Λ	S	9	•	•	R	d	Н	1	Ţ	•	•	Λ	•	3	I	V	•	٨	•	•	٨
L3(B)	_	۸	S	9	•	•	К	•	Н	7	1	•	•	٨	Н	Ę	I	•	•	•	•	•	•
L9(A)		•		•	•	^	•	•	•	•	•	•	•	٠	•	•	•	A	٠	•	^		•
L8(A)	•	•	•	•	•	٨	٠	•	•	•	•	•	N	•	٠	•	•	•	•	•	•	Ь	•
(A)97	•	•	•	•	•	•	•	•	•	•	•	•	•	•	٠	•	•	•	•	•	•	•	•
L2(A)		•	•	•	•	•	•	•	•	٠	•	•	٠	•	•	. •	•	•	×	٠	•	•	•
L1* (A)		•	•	•		^	•	•	·	•	•	•	•	•	•	•	•	•	•	•	٨	•	•
Cons-p9	u	H	Ţ	Y.	^	A	I	S	O	L	A	S	S	٧	λ	У	۸	>	۸	I	7	ð	Y
L fragment	384	386	388	390	391	392	394	405	434	438	444	450	458	466	474	476	496	524	536	580	622	673	783
	F2	!																					/d

FIG. 7B

Cons-F	9	W	×	\ \ \ \	A	A,D	ط	A	V,I	H,O		Ð	S	ليا	A	A	¥	}	> -	├ ─	7
Cons-D		•	•	•		۵	•			0,H		•	•		•	•	•		•		•
L4(C)	•	•		•		•		•		•					•	•	•				
L10(B)		•		•		Ω			•	•			•		^	•	•	S			•
(B) _* (7)	S		8			٥		Ь	I	I	•		•	•	•	•	•	. •	•		•
L3(B)	•	I		•		Q	•	•	I	エ			•	•	•	•	•	S	•		•
(A)61	•	•	•				•	•	_	I	•		•		•	•	•		•	•	•
L8(A)	•	•	•		•	•	S	•		•		•	•		•	•	•		•	•	•
L6(A)	•	•	•		^		•	•	Ι	•	•		1		•	•	•		•		•
L2(A)	•	•	•	П		•	•			•			•	S	•	•	•	•	•		Ь
L1* (A)	•	•	•	_			•		•	•	Χ	~	•		•	٨	Z	•	•	٧	•
Cons-p9	9	W	¥	^	A	A	Ь	A	^	Ö	·	9	S	ببا	A	A	¥		У	<u></u>	
L fragment	820	857	927	934	937	978	1028	1031	1043	1067	1097	1188	1215	1223	1226	1339	1399	1503	1528	1535	1662
:	NS2			•		·	NS3							•				•			NS4A

FIG. 7C

	<u> </u>	,	,		_				,	_					_	_			,		_	
Cons-F	ϫ	N,T	S	≥	~	∀ '⊥			E,D	>	1,0	>	z	A	O		A		S	S	۵	S
Cons-D	•	z	•	•		A,T	•		E,D	•	۱,۵	•	•	•	•				•			
(C)		z	۵.	•	~							T		•		•		98		9	ပ	
L10(B)		z				A			Ω	•	0			•				S4		9		
(B) _* (C)	•	•	•	•	•	A	•		Q	•	Ö	•	S	•	•	•	•	S10	S	S		•
L3(B)	•	Z	•	I		A	۵.	ட	O	Ι	Q	•	•	•	•	•	^	S8	S	9	•	•
L9(A)	•	•	•	٨		•	•	•	•	•	•			•	•	•		S7	•	•	•	•
L8(A)	•	•	•	•	•	•	•	•	•	٠	•	•	•	•	•	•	•	S3	•	•	•	F
L6(A)	•	N	•	•	•	•	•	•	•	•	•	•	•	•	•	•	•	S2	•	•	•	Ŀ
L2(A)	Р	•	•	•	•	•	•	•	•	•	•	•	•	•	R	٨	•	29	•	•	•	4
L1* (A)	•	•	•	•	•	٠	•		٠	•	•	•	•	•	•	•	•	S5	•	•	•	•
Cons-p9	У	Н	S	M	Х	Ţ		7	Ε	٨	٦	Y	N	А	Ö	I	A		9	S	Ō	S
L fragment	1753	1805	1949	2105	2136	2146	2226	2259	2262	2334	2371	2385	2692	2757	2785	2824	2861	S fragment	2968	2975	2978	2999
	NS4B			NS5A									NS5B									

FIG. 7D

	L1 (A)	L2 (A)	L6 (A)	L8 (A)	(A) (L9	(B)	(8)	L10 (B)	L4 (C)	HC-J4/91HC-J4/83	HC-J4/83
/		0.56	09.0	0.36	0.33	1.50	1.53	1.46	0.95	0.83	1.79
0	0.59		0.55	0.35	0.50	1.49	1.51	1.45	0.98	0.82	1.77
0	0.52	0.42		0.31	0.55	1.33	1.38	1.29	08.0	0.68	1.58
0	0.42	0.38	0.31		0.31	1.32	1.34	1.28	0.79	0.65	1.62
0	0.35	0.52	0.45	0.35		1.42	1.42	1.38	0.91	0.75	1.66
1	1.47	1.43	1.15	1.33	1.36		0.61	0.30	1.43	06.0	1.51
	1.36	1.33	1.05	1.22	1.22	99.0		0.57	1.47	0.95	1.54
1	.36	1.33	0.59	1.22	1.26	0.31	0.56		1.37	0.85	1.42
0	0.77	0.80	0.59	69.0	1.26	1.12	1.08	1.01		0.76	1.73
٦	0.94	0.91	0.63	08.0	0.87	0.77	0.73	99.0	0.52		1.22
-	1.96	1.89	1.68	1.85	1.82	1.75	1.61	1.61	1.71	1.40	
ĺ											7

FIG. 8

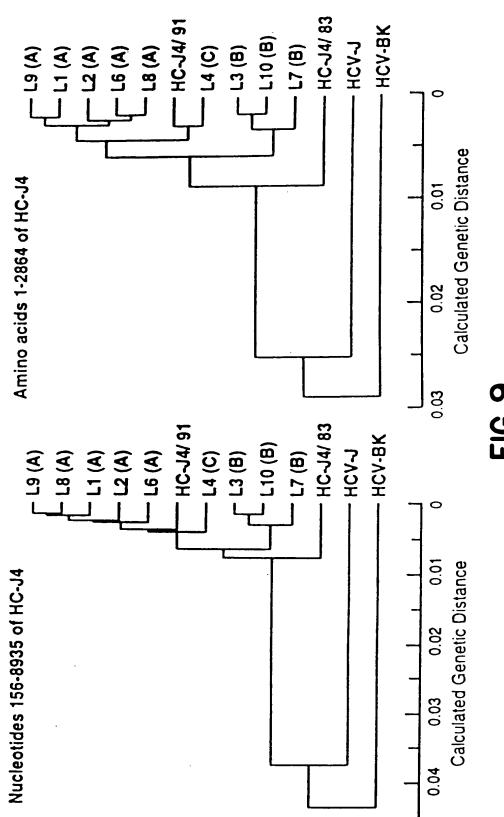


FIG. 9

	379		413	468		486
HC-J4L6 (A)	: AGVDG	3 ETHTTGRVAGHTTSGFTSLFSSGAS	QKIQL	GWGPIT	YTKPNSS	DQRPYC
HC-J4L2 (A)	•		•	•		•
HC-J4/91-20	•	•	•	•		•
	•		•	•	•	•
HC-J4L8 (A)	•		•	•	•	•
	•			•	•	•
HC-J4/91-21	•	>		•	 G	•
HC-J4L4 (C)	•		•	•	ED:	•
HC-J4/91-23	••	•	•	•	ю :	•
HC-J4/91-22	•		•	A		•
HC-J4L7 (B)	•	T.Y.	:	•	 	•
0	T		•	•		•
HC-J4L3 (B)	H	T.Y.	•	•	H.E	•
HC-J4/91-26	H	T.Y.	•	•	G.D.L	•
HC-J4/91-25	•	A.Y.S	•	•	ы	•
HC-J4/91-24	•	છ	•	•	. Е Р	•
HC-J4/91	•	A.Y.S	•	•	EP	:
HC-J4/91-27	•	K.Y.S.GA.SRP	.R	•	.ESG.R	•
HC-J4/83	•	Y.S	. R	•	E.D.P	•
		HVR1 FIG. 10	<u>o</u>		HVR2	

Series and the series of the s

5' Untranslated Region		06
HC-J4 :GCCAGCCCC GATIGGGGGC GACACTCCAC CATAGATCAC TCCCCTGTGA GG pCV-J4L6S: TGA	GGAACTACTG	TCTTCACGCA GAAAGCGTCT AGCCATGGCG
91 HC-J4 :TTAGTATGAG TGTCGTGCAG CCTCCAGGAC CCCCCTCCC GGGAGAGCCA TAC pCV-J4L6S:	TAGTGGTCTG	CGGAACCGGT GAGTACACCG GAATTGCCAG
181 HC-J4 :GACGACCGGG TCCTTTCTTG GATCAACCCG CTCAATGCCT GGAGATTTGG GC pCV-J4L6S:	SATTIGG GCGIGCCCCC GCGAGACIGC	GCGAGACTGC TAGCCGAGTA GTGTTGGGTC
271 HC-J4 :GCGAAAGGCC TIGIGGTACT GCCTGATAGG GTGCTTGCGA GTGCCCCGGG AGGTCTCGTA pCV-J4L6S:	CCCCGGG AGGTCTCGTA	341 GACCGTGCAC C
3' Untranslated Region		
3' variable region poly U-U	poly U-UC region	3' variable region
9372 HC-J4 :IGAACGGGGA GCTAACCACT CCAGGCCAAT AGGCCTTC CTG poly pCV-J4L6S:poly pCV-H77C :G.C.TCTAA.TT poly bCV-H77C :G.TT.G .GA		9513 GGTGGCT CCATCTTAG AAT
3' conserved region (Cont.)	on (Cont.)	
GAAAGGTCCG TGAGCCGCAT	GACTGCAGAG AGTGCTGATA	9595 CIGGCCTCTC IGCAGAICAI GI
FIG. 11		

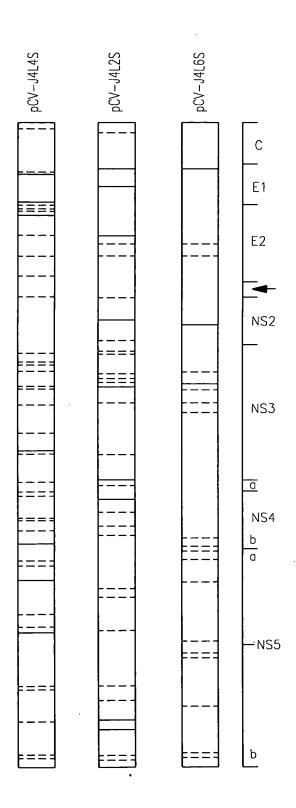


FIG. 12

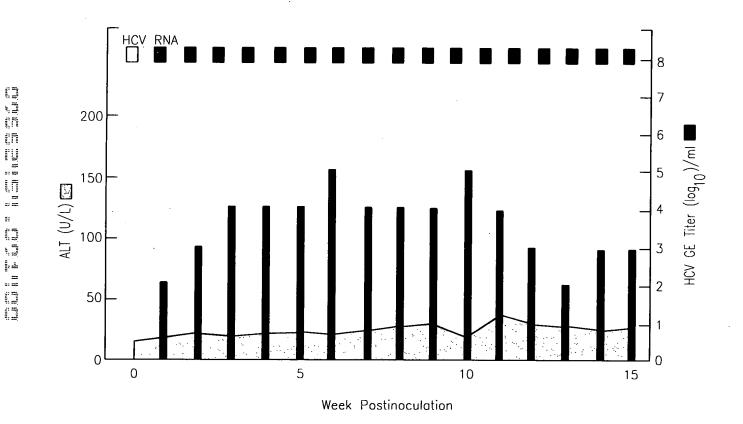


FIG. 13

					
10	20	30	40	50	
			1234567890		
			CATGAATCAC		50
			AGCCATGGCG		100
TGTCGTGCAG	CCTCCAGGAC	CCCCCCTCCC	GGGAGAGCCA	TAGIGGICIG	150
CGGAACCGGT	GAGTACACCG	GAATTGCCAG	GACGACCGGG	TCCTTTCTTG	200
GATCAACCCG	CICAATGCCT	GGAGATTTGG	GCGIGCCCCC	GCGAGACTGC	250
TAGCCGAGIA	GIGIIGGGIC	GCGAAAGGCC	TIGIGGIACT	GCCTGATAGG	300
GIGCTIGCGA	GIGCCCCCGGG	AGGICICGIA	GACCGTGCAC	CATGAGCACG	350
AATCCTAAAC	CTCAAAGAAA	AACCAAACGT	AACACCAACC	GCCGCCCACA	400
GGACGTCAAG	TTCCCGGGGGG	GIGGICAGAT	CGTTGGTGGA	GITTACCIGT	450
TGCCGCGCAG	GGGCCCCAGG	TIGGGIGIGC	GCGCGACTAG	GAAGGCTTCC	500
GAGCGGTCGC	AACCTCGTGG	AAGGCGACAA	CCTATCCCAA	AGGCTCGCCG	550
ACCCGAGGGC	AGGGCCTGGG	CTCAGCCCGG	GIACCCITGG	CCCCTCTATG	600
GCAATGAGGG	CCIGGGGIGG	GCAGGATGGC	TCCTGTCACC	CCCCCCCCTCC	650
CGGCCTAGIT	GGGGCCCCAC	GGACCCCCGG	CGTAGGTCGC	GTAACTTGGG	700
TAAGGICATC	GATACCCTTA	CATGCGGCTT	CGCCGATCTC	ATGGGGTACA	750
TTCCGCTCGT	aggagaaca	CTAGGGGGGG	CTGCCAGGGC	CTTGGCACAC	800
GGIGICCGGG	TTCTGGAGGA	CGGCGTGAAC	TATGCAACAG	GGAACTTGCC	850
CCGTTCCTCT	TICICIATCT	TCCTCTTGGC	TCIGCIGICC	TGTTTGACCA	900
TCCCAGCTTC	CGCTTATGAA	GIGCGCAACG	TGTCCCGGGAT	ATACCATGTC	950
ACGAACGACT	GCTCCAACTC	AAGCATIGIG	TATGAGGCAG	CCCACCICAT	1000
CATGCATACT	CCCCGGGTGCG	TECCCIGIGI	TCAGGAGGGT	AACAGCTCCC	1050
GITGCIGGGI	AGCGCTCACT	CCCACGCTCG	CGGCCAGGAA	TGCCAGCGTC	1100
CCCACTACGA	CAATACGACG	CCACGICGAC	TICCICCITG	GGACGGCTGC	1150
TTTCTGCTCC	GCTATGTACG	TGGGGGATCT	CIGCGGATCT	ATTTTCCTCG	1200
TCTCCCAGCT	GTTCACCTTC	TCGCCTCGCC	GGCATGAGAC	AGTGCAGGAC	1250
TGCAACTGCT	CAATCTATCC	CGGCCATGTA	TCAGGTCACC	GCATGGCTTG	1300
GGATATGATG	ATGAACTGGT	CACCTACAAC	ACCCCTAGTG	GIGICGCAGI	1350
TGCTCCGGAT	CCCACAAGCT	GICGIGGACA	TGGTGGCGGG	GGCCCACTGG	1400
GGAGTCCTGG	CCCCCTTCC	CIACIATICC	ATGGTAGGGA	ACTGGGCTAA	1450
GGTTCTGATT	GIGGCGCTAC	TCTTTGCCCGG	CGTTGACGGG	GAGACCCACA	1500
CGACGGGGAG	GGIGGCCGGC	CACACCACCT	CCCCCTTCAC	GICCCTTTTC	1550
TCATCTGGGG	CGICICAGAA	AATCCAGCTT	GIGAATACCA	ACGGCAGCTG	1600
GCACATCAAC	AGGACTGCCC	TAAATTGCAA	TGACTCCCTC	CAAACTGGGT	1650
	· · · · · · · · · · · · · · · · · · ·		TCAACICGIC		1700
GAGCGCATGG	CCAGCIGCCG	CCCCATIGAC	TEGTTCECCC	AGGGGTGGGG	1750
CCCCATCACC	TATACTAAGC	CTAACAGCTC	GGATCAGAGG	CCTTATIGCT	1800
GGCATTACGC	GCCTCGACCG	TGIGGIGICG	TACCCGCGTC	GCAGGIGIGI	1850
GGTCCAGTGT	ATTGTTTCAC	CCCAAGCCCT	GIIGIGGIGG	GGACCACCGA	1900

FIG. 14A

10	20	30	40	50	
1234567890 1234	<u> 1567890 12</u>	<u>34567890</u>	1234567890	1234567890	·
TOGITOCOGIT GIO	CTACGT AT	AGCTGGGG	GGAGAATGAG	ACAGACGIGA	1950
TGCTCCTCAA CAAC	CACGCGT CC	GCCACAAG	GCAACIGGIT	CGGCTGTACA	2000
TGGATGAATA GTAC	CIGGGIT CA	CTAAGACG	TGCGGAGGIC	CCCCGIGIAA	2050
CATCGGGGGG GTCC	GIAACC GC	ACCITGAT	CIGCCCCACG	GACTGCTTCC	2100
GGAAGCACCC CGAC	GCTACT TA	CACAAAAT	GIGGCIGGGG	GCCCIGGIIG	2150
ACACCTAGGT GCC	PAGTAGA CI	ACCCATAC	AGGCTTTGGC	ACIACCCCIG	2200
CACICICAAT TIT	ICCAICT II	AAGGTTAG	CATCIATGIG	GGGGGGGGIGG	2250
AGCACAGGCT CAA	IGCCGCA TG	CAATTOGA	CTCCACCACA	GCGCTGTAAC	2300
TTGGAGGACA GGG	ATAGGIC AG	AACTCAGC	CCCCTCCTCC	TGICTACAAC	2350
AGAGTGGCAG ATA	CIGCCCT GI	CTTTCAC	CACCCTACCG	GCTTTATCCA	2400
CIGGITIGAT CCA	ICICCAT CA	GAACATCG	TOGACGTOCA	ATACCIGIAC	2450
GGTGTAGGGT CAG	CETTET CI	CCTTTCCA	ATCAAATGGG	AGTACATCCT	2500
GITGCITTIC CIT	CTCCTGG CA	GACGCGCG	CCICICICCC	TECTTETEGA	2550
TGATGCTGCT GAT	AGCCCAG GC	TGAGGCCG	CCTTAGAGAA	CITEGIGGIC	2600
CTCAATGCGG CGT	CCCTGGC CC	GAGCGCAT	GGTATTCTCT	CCTTTCTTGT	2650
GITCITCIGC GCC	GCCTGGT AC	ATTAAGGG	CAGGCTGGCT	CCTGGGGGGG	2700
CGTATGCTTT TTA	TGGCGIA TG	ECCOCCTCC	TCCTGCTCCT	ACTOGCGTTA	2750
CCACCACGAG CIT	ACGCCTT GC	EACCGGGAG	ATGGCTGCAT	CGTGCGGGGG	2800
TGCGGTTCTT GTA	GGICIGG TA	ATTCTTGAC	CTTGTCACCA	TACTACAAAG	2850
TGTTTCTCAC TAG	GCTCATA TO	GIGGITAC	AATACTTTAT	CACCAGAGCC	2900
GAGGCGCACA TGC	AAGIGIG GO	TOCCOCC	CTCAACGITC	GGGGAGGCCG	2950
CGATGCCATC ATC	CICCICA CO	FIGICCET	TCATCCAGAG	TTAATTTTTG	3000
ACATCACCAA ACT	CCIGCIC GO	CATACICG	GCCCGCTCAT	GGTGCTCCAG	3050
GCTGGCATAA CGA	GAGTGCC GT	PACTICGIG	CGCGCTCAAG	GCCTCATTCG	3100
TGCATGCATG TTA	GTGCGAA AA	AGTCGCCGG	GGGTCATTAT	GTCCAAATGG	3150
TCTTCATGAA GCT	GGGCGCC C.1	IGACAGGIA	CGTACGTTTA	TAACCATCTT	3200
ACCCCACTGC GGG	ACTGGGC CO	CACGCGGGC	CTACGAGACC	TICCCGICCC	3250
GGTAGAGCCC GTC	GICTICT CO	CCCATCGA	GACCAAGGIC	ATCACCTGGG	3300
GAGCAGACAC CGC	TECGIGI C	GGACATCA	TCTTGGGTCT	ACCCGICICC	3350
GCCCGAAGGG GGA	AGGAGAT A	TTTTTCCCA	CCGGCTGATA	GICICGAAGG	3400
GCAAGGGTGG CGA	CICCIIG C	SCCCATCAC	GGCCTACTCC	CAACAAACGC	3450
GGGGGTACT TGG	TIGCATC A	ICACTAGCC	TCACAGGCCG	GCACAAGAAC	3500
CAGGICGAAG GGG	EAGGITCA A	GIGGITICI	ACCGCAACAC	AATCTTTCCT	3550
GGCGACCTGC ATO					
CGAAGACCCT ACC					
GTAGACCTGG ACC	CICCICCC C	TGGCAGGCG	CCCCCCCGGGG	CGCGCTCCAT	
GACACCATGC AGO	MGTGGCA G	CICGGACCI	TIACTIGGIC	ACGAGACATG	
CTGATGICAT TO	COCTOCOC C	GGCGAGGCG	ACAGCAGGGG	AAGICIACIC	3800

FIG. 14B

						
10	20	30	40	50		
			1234567890			_
			TCCTCGGGTG		3850	
			CCCGCCTCCT		3900	
			CCGTTGAGIC	· · · 	3950	
			TCAACCCCCC		4000	
			TCCTACTGGC	·	4050	
GCACCAAAGT	GCCGGCIGCG	TATGCAGCCC	AAGGGTACAA	GGIGCICGIC	4100	
CTGAACCCGT	CCCLLCCCCC	CACCTTAGGG	TITIGGGGGGGT	ATATGTCCAA	4150	
GGCACACGGT	ATCGACCCTA	ACATCAGAAC	TGGGGTAAGG	ACCATTACCA	4200	
CGGGCGGCIC	CATTACGIAC	TCCACCTATG	GCAAGTTCCT	TGCCGACGGT	4250	
CCCIGITCIG	GGGGGCCTA	TGACATCATA	ATATGTGATG	AGTGCCACTC	4300	
AACIGACICG	ACTACCATCT	TGGGCATCGG	CACAGTOCTG	GACCAAGCGG	4350	
AGACGGCTGG	AGCGCGGCTC	GICGICCICG	CCACCGCTAC	ACCTCCGGGA	44 00	
TCGGTTACCG	TGCCACACCC	CAATATCGAG	GAAATAGGCC	TGTCCAACAA	4450	
TGGAGAGATC	CCCTTCTATG	GCAAAGCCAT	CCCCATTGAG	GCCATCAAGG	4500	
GGGGGAGGCA	TCTCATTTTC	TGCCATTCCA	AGAAGAAATG	TGACGAGCTC	4550	
GCCGCAAAGC	TGACAGGCCT	CGGACTGAAC	GCTGTAGCAT	ATTACCGGGG	4600	
CCTTGATGTG	TCCGTCATAC	CGCCTATCGG	AGACGICGIT	GTCGTGGCAA	4650	
CAGACGCTCT	AATGACGGGT	TTCACCGGCG	ATTTTGACTC	AGTGATCGAC	4700	
TGCAATACAT	GIGICACCCA	GACAGTCGAC	TICAGCTIGG	ATCCCACCIT	4750	
CACCATTGAG	ACGACGACCG	TGCCCCAAGA	CCCCCTCTCCC	CGCTCGCAAC	4800	
CCCCACCTAC	AACTGGCAGG	CCTACCACTC	GCATCTACAG	GITIGIGACT	4850	
CCAGGAGAAC	GGCCCICGGG	CATGITCGAT	TCITCCGTCC	TGIGIGAGIG	4900	
CTATGACGCG	CCCTCTCCTT	GGTATGAGCT	CACGCCCCCT	GAGACCTCCG	4950	
TTAGGTTGCG	GGCTTACCTA	AATACACCAG	GGTTGCCCGT	CIGCCAGGAC	5000	
CATCTGGAGT	TCTGGGAGAG	CGTCTTCACA	GGCCTCACCC	ACATAGATGC	5050	
CCACTTCCTG	TCCCAGACTA	AACAGGCAGG	AGACAACTTT	CCTTACCTCG	5100	
TGGCATATCA	AGCTACAGTG	TGCGCCAGGG	CTCAAGCTCC	ACCTCCATCG	5150	
TGGGACCAAA	TGTGGAAGTG	TCTCATACGG	CTGAAACCTA	CACTGCACGG	5200	
GCCAACACCC	CIGCIGIATA	GGCTAGGAGC	CGTCCAAAAT	GAGGICATCC	5250	
TCACACACCC	CATAACTAAA	TACATCATGG	CATGCATGIC	GGCTGACCTG	5300	
GAGGICGICA	CTAGCACCTG	GGTGCTGGTA	GGCGGAGTCC	TIGCAGCITT	5350	
GGCCGCATAC	TGCCTGACGA	CAGGCAGIGI	GGICATIGIG	GGCAGGATCA	5400	
TCTTGTCCCG	GAAGCCAGCT	GICGIICCCG	ACAGGGAAGT	CCTCTACCAG	5450	
GAGTTCGATG	AGATOGAAGA	GIGIGCCICA	CAACTTCCTT	ACATCGAGCA	5500	
GGGAATGCAG	CTCGCCGAGC	AATTCAAGCA	AAAGGCGCTC	CCCTTCTTCC	5550	
AAACGGCCAC	CAAGCAAGCG	GAGGCTGCTG	CTCCCGTGGT	GGAGTCCAAG	5600	
TGGCGAGCCC	TTGAGACCTT	CTGGGCGAAG	CACATGIGGA	ATTTCATCAG	5650	
CGGAATACAG	TACCTAGCAG	GCTTATCCAC	TCTGCCTGGA	AACCCCCGCGA	5700	

FIG. 14C

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
TAGCATCATT	GATGGCATTT	ACAGCTTCTA	TCACTAGCCC	GCTCACCACC	5750
CAAAACACCC	TCCIGITIAA	CATCTTGGGG	GGATGGGTGG	CTGCCCAACT	5800
CCCTCCTCCC	AGCGCIGCGI	CAGCTTTCGT	GGGGGGGGGC	ATCCCCCCGAG	5850
CCCCTCTTCC	CAGCATAGGC	CTTGGGAAGG	TGCTCGTGGA	CATCITGGCG	5900
GGCTATGGGG	CAGGGGTAGC	CGGCGCACTC	GIGGCCITIA	AGGTCATGAG	5950
CCCCCACCIG	CCCTCCACCG	AGGACCTGGT	CAACITACIC	CCIGCCATCC	6000
TCICICCICG	TGCCCTGGTC	GICCGCGGICC	TGTGCGCAGC	AATACTGCGT	6050
CGGCACGIGG	GCCCCGGGAGA	GGGGGCIGIG	CAGIGGATGA	ACCCCCTCAT	6100
AGOGITOGCT	TCGCCGGGTA	ACCACGICIC	CCCTACGCAC	TATGIGCCIG	6150
AGAGCGACGC	TGCAGCACGT	GICACICAGA	TCCTCTCTAG	CCTTACCATC	6200
ACTCAACTGC	TGAAGCGGCT	CCACCAGIGG	ATTAATGAGG	ACIGCICIAC	6250
GCCATGCTCC	GCCICCIGCC	TAAGGGATGT	TICCCATICG	ATATGCACGG	6300
TGTTGACTGA	CTTCAAGACC	TGGCTCCAGT	CCAAACTCCT	GCCGCGGTTA	6350
CCGGGAGTCC	CTTTCCTGTC	ATGCCAACGC	GGGTACAAGG	GAGICIGGCG	6400
GGGGGACGGC	ATCATGCAAA	CCACCTGCCC	ATGCGGAGCA	CAGATCGCCG	6450
GACATGICAA	AAACGGTTCC	ATGAGGATCG	TAGGGCCTAG	AACCTGCAGC	6500
AACACGIGGC	ACGGAACGIT	CCCCATCAAC	CCATACACCA	CCCCACCTTG	6550
CACACCCICC	CCGGCGCCCA	ACTATTCCAG	GGCGCTATGG	CCCCTC	6600
CTGAGGAGTA	CGTGGAGGTT	ACGCGTGTGG	GGGATTTCCA	CTACGIGACG	6650
GGCATGACCA	CIGACAACGT	AAAGTGCCCA	TGCCAGGTTC	CGGCCCCGA	6700
ATTCTTCACG	GAGGIGGATG	GAGIGCGGIT	GCACAGGTAC	GCTCCGGCGT	6750
GCAAACCICT	TCTACGGGAG	GACGICACGI	TCCAGGTCGG	GCTCAACCAA	6800
TACITOGICG	GGTCGCAGCT	CCCATGCGAG	CCCGAACCGG	ACGTAACAGT	6850
GCTTACTTCC	ATGCTCACCG	ATCCCTCCCA	CATTACAGCA	GAGACGGCTA	6900
AGCGTAGGCT	GGCTAGAGGG	TCTCCCCCCT	CTTTAGCCAG	CICATCAGCT	6950
AGCCAGTIGT	CIGCGCCTIC	TTTGAAGGCG	ACATGCACTA	CCCACCATGA	7000
CTCCCCGGAC	GCTGACCTCA	TCGAGGCCAA	CCICITGIGG	CGGCAGGAGA	7050
		GIGGAGICAG			7100
		CGCGGAGGGG			7150
· -		AATCCAGGAA			7200
		AATCCTCCAC			7250
		GGTACACGGA		· · · · - - · ·	7300
		CACGGAGAAA			7350
•		TTGGCGGAGC			7400
		TGATAGCGGC			7450
		ACAAAGGATC			7500
		GAGCCGGGGG			7550
TCTTGGTCTA	. CCGTGAGTGA	GGAGGCTAGT	GAGGATGTCG	TCTCCTCCTC	7600

FIG. 14D

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
AATGICCIAT	ACGIGGACAG	GCGCCCTGAT	CACGCCATGC	GCTGCCGGAGG	7650
AAAGIAAGCT	GCCCATCAAC	COGTTGAGCA	ACICITICCT	GOGTCACCAC	7700
AACATGGTCT	ACGCCACAAC	ATCCCGCAGC	GCAAGCCTCC	GGCAGAAGAA	7750
GGICACCITT	GACAGATTGC	AAGICCIGGA	TGATCATTAC	CCCCACCTAC	7800
TCAAGGAGAT	GAAGGCGAAG	GCGTCCACAG	TTAAGGCTAA	GCTTCTATCT	7850
ATAGAGGAGG	CCTGCAAGCT	GACGCCCCCA	CATTOGGCCA	AATCCAAATT	7900
TGGCTATGGG	GCAAAGGACG	TCCGGAACCT	ATCCAGCAGG	GCCGITAACC	7950
ACATCCCCTC	CGTGTGGGAG	GACTTCCTCC	AAGACACTGA	AACACCAATT	8000
GACACCACCA	TCATGGCAAA	AAGTGAGGTT	TICIGOGICC	AACCAGAGAA	8050
GGGAGGCCGC	AAGCCAGCTC	GCCTTATCGT	ATTCCCAGAC	CIGGGAGITC	8100
GIGIATCCGA	GAAGATGGCC	CTTTACGACG	TEGTCTCCAC	CCTTCCTCAG	8150
GCCGTGATGG	GCTCCTCATA	CGGATTTCAA	TACTCCCCCA	AGCAGCGGGT	8200
CGAGTTCCTG	GIGAATACCT	GGAAATCAAA	GAAATGCCCT	ATGGGCTTCT	8250
CATATGACAC	CCCCTGTTTT	GACTCAACGG	TCACTGAGAG	TGACATTCGT	8300
GIIGAGGAGI	CAATTTACCA	ATGITGIGAC	TTGGCCCCCG	AGGCCAGACA	8350
GGCCATAAGG	TOGCTCACAG	AGCGGCTTTA	CATCGGGGGT	CCCCTGACTA	8400
ACTCAAAAGG	GCAGAACTGC	GCTTATCGCC	GGIGCCGCGC	AAGTGGCGTG	8450
CIGACGACTA	GCTGCGGTAA	TACCCTCACA	TGITACTIGA	AGGCCACTGC	8500
AGCCTGTCGA	GCTGCAAAGC	TCCAGGACTG	CACGATGCTC	GTGAACGGAG	8550
ACGACCTTGT	CGITATCIGI	GAAAGCGCGG	GAACCCAGGA	GGATGCGGCG	8600
GCCCTACGAG	CCTTCACGGA	GCTATGACT	AGGTATTCCG	CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	8650
GGATCCGCCC	CAACCAGAAT	ACGACCTGGA	GCTGATAACA	TCATGTTCCT	8700
CCAATGIGIC	AGTOGOGCAC	GATGCATCIG	GCAAAAGGGT	ATACTACCTC	8750
ACCCGTGACC	CCACCACCCC	CCTTGCACGG	GCIGCGIGGG	AGACAGCTAG	8800
ACACACTCCA	ATCAACTCTT	GUCTAGGCAA	TATCATCATG	TATGCGCCCA	8850
CCCTATGGGC	AAGGATGATT	CIGATGACTC	ACTITITCIC	CATCCTTCTA	8900
GCTCAAGAGC	AACTIGAAAA	AGCCCTGGAT	TGTCAGATCT	ACGCCCTTC	8950
CTACTCCATT	GAGCCACTTG	ACCTACCTCA	GATCATTGAA	CCACTCCATG	9000
GICTTAGCGC	ATTTACACTC	CACAGITACT	CICCAGGIGA	GATCAATAGG	9050
GIGGCTICAT	GCCTCAGGAA	ACTTGGGGTA	CCACCCTTGC	GAACCTGGAG	9100
ACATCGGGCC	AGAAGIGICC	GCGCTAAGCT	ACTGTCCCAG	GGGGGGAGGG	9150
CCGCCACTIG	TGGCAGATAC	CICTTIAACT	GGGCAGTAAG	GACCAAGCTT	9200
		CGCGTCCCAG			9250
		GAGACATATA			9300
		TGCCTACTCC			9350
		ATGAACGGGG			9400
		TTTTTTTTT			9450
TITCITICCT	TICCTICTIT	TTTTCCTTTC	TTTTTCCCTT	CTTTAATGGT	9500

FIG. 14E

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	
GGCTCCATCT	TAGCCCTAGT	CACGGCTAGC	TGTGAAAGGT	CCGTGAGCCG	9550
CATGACTGCA	GAGAGIGCIG	ATACTGGCCT	CTCTGCAGAT	CAIGT	9595

FIG. 14F

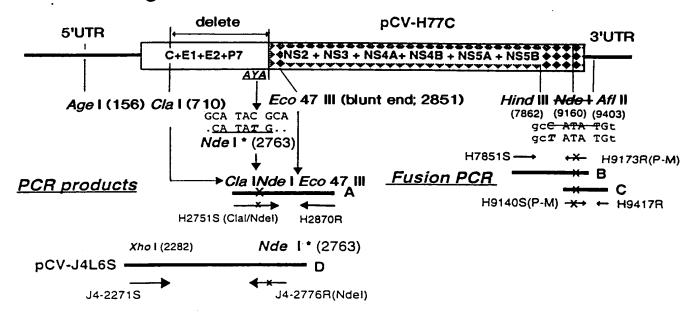
10	20	30	40	50	
			1234567890		
			VGGVYLLPRR		50
		-	YPWPLYGNEG	_	100
			ADLMGYIPLV	· ·	150
LAHGVRVLED	GVNYATGVLP	GCSFSIFLLA	LLSCLTIPAS	AYEVRNVSGI	200
YHVINDCSNS	SIVYEAADVI	MHITGCVPCV	QEGNSSROW	ALTPILAARN	250
ASVPITITIRR	HVDLLVGTAA	FCSAMYVGDL	CGSIFLVSQL	FIFSPRRHET	300
VQDCNCSIYP	CHVSCHRMAW	DMMNWSPTT	ALVVSQLLRI	PQAVVDMVAG	350
AHWGVLAGLA	YYSMVGVWAK	VLIVALLFAG	VDGETHITGR	VAGHITSGFT	400
SLFSSGASQK	IQLVNINGSW	HINRIALNON	DSLQIGFFAA	LFYAHKFNSS	450
GCPERMASCR	PIDWFAQGWG	PITYTKPNSS	DQRPYCWHYA	PRPCGVVPAS	500
QVCGPVYCFT	PSPVVVGTTD	RSGVPTYSWG	ENETDVMLLN	NIRPPQGNWF	550
CCIWMNSTGF	TKTCGGPPCN	IGGVGNRILI	CPIDCFRKHP	EATYTKCGSG	600
PWLTPRCLVD	YPYRLWHYPC	TLNFSIFKVR	MYVGGVEHRL	NAACNWIRGE	650
RCNLEDRDRS	ELSPLLLSTT	EWQILPCAFT	TLPALSTGLI	HLHQNIVDVQ	700
YLYGVGSAFV	SFAIKWEYIL	LLFLLLADAR	VCACLWMMLL	IAQAEAALEN	750
LVVLNAASVA	GAHGILSFLV	FFCAAWYIKG	RLAPGAAYAF	YGWPLLLLL	800
LALPPRAYAL	DREMAASCGG	AVLVGLVFLT	LSPYYKVFLT	RLIWWLQYFI	850
TRAEAHMQVW	VPPLNVRGGR	DAIILLTCAV	HPELIFDITK	LLLATIGPIM	900
VLQAGITRVP	YFVRAQGLIR	${\tt ACMLVRKVAG}$	$\hbox{CHYVQMVFMK}$	LGALIGIYVY	950
			TKVITWGADT	· · · · · · · · · · · · · · · · · · ·	1000
PVSARRCKEI	FLGPADSLEG	QGWRLLAPIT	AYSQQTRGVL	CCITTSLITCER	1050
DKNQVEGEVQ	VVSTATQSFL	ATCINGVCWT	VYHGAGSKTL	AGPKGPITQM	1100
YINVDLDLVG	WQAPPGARSM	TPCSCGSSDL	YLVTRHADVI	PVRRRGDSRG	1150
SLLSPRPVSY	LKGSSGGPLL	CPSGHVVGVF	RAAVCTRGVA	KAVDFIPVES	1200
METTIMRSPVF	TDNSTPPAVP	QTFQVAHLHA	PIGSGKSTKV	PAAYAAQGYK	1250
VLVLNPSVAA	TLGFGAYMSK	AHGIDPNIRT	GVRTTTTGGS	ITYSTYGKFL	1300
ADGGCSGGAY	DIIICDECHS	TESTITLGIG	TVLDQAETAG	ARLVVLATAT	1350
PPGSVIVPHP	NIEEIGLSNN	GEIPFYGKAI	PIEAIKGGRH	LIFCHSKKKC	1400
DELAAKLIGL	GLNAVAYYRG	LDVSVIPPIG	DVVVVATDAL	MIGFIGDFDS	1450
· -			AVSRSQRRGR		1500
			TPAETSVRLR		1550
			DNFPYLVAYQ		1600
PPSWDQMWKC	LIRLKPTLHG	PTPLLYRLGA	VQNEVILTHP	ITKYIMACMS	1650
			VIVGRIILSG		1700
			KALGLLQTAT		1750
			LPGNPAIASL		1800
			GAGIAGAAVG		1850
ILAGYGAGVA	GALVAFKVMS	GEVPSTEDLV	NLLPAILSPG	ALVVGVVCAA	1900

FIG. 14G

10	20	30	40	50	
1234567890	1234567890	1234567890	1234567890	1234567890	·
ILRRHVGPGE	GAVQWMNRLI	AFASRGNHVS	PIHYVPESDA	AARVIQILSS	1950
LTTTQLLKRL	HOWINEDCST	PCSGSWLRDV	WDWICIVLID	FKIWLQSKLL	2000
PRLPGVPFLS	CORGYKGVWR	GDGIMQITICP	CGAQIAGHVK	NGSMRIVGPR	2050
TCSNIWHGIF	PINAYTIGPC	TPSPAPNYSR	ALWRVAAEEY	VEVIRVGDFH	2100
YVIGMITDIV	KCPCQVPAPE	FFTEVDGVRL	HRYAPACKPL	LREDVITQVG	2150
LNQYLVGSQL	PCEPEPDVIV	LISMLIDPSH	ITAETAKRRL	ARGSPPSLAS	2200
SSASQLSAPS	LKATCTTHHD	SPDADLIEAN	LLWRQEMGGN	TTRVESENKV	2250
VILDSFEPLH	AEGDEREISV	AAETLRKSRK	FPSALPIWAR	PDYNPPLLES	2300
WKDPDYVPPV	VHGCPLPPIK	APPIPPPRRK	RIVVLIESNV	SSALAELATK	2350
TFGSSGSSAV	DSGTATALPD	LASDDGDKGS	DVESYSSMPP	LEGEPGDPDL	2400
SDGSWSTVSE	EASEDVVCCS	MSYIWICALI	TPCAAEESKL	PINPLSNSLL	2450
RHHNMVYATT	SRSASLRQKK	VIFDRLQVLD	DHYRDVLKEM	KAKASTVKAK	2500
LLSIEEACKL	TPPHSAKSKF	GYGAKDVRNL	SSRAVNHIRS	WEDLLEDIE	2550
TPIDITIMAK	SEVFCVQPEK	GCRKPARLIV	FPDLGVRVCE	KMALYDVVST	2600
LPQAVMGSSY	GFQYSPKQRV	EFLVNIWKSK	KCPMGFSYDT	RCFDSIVIES	2650
DIRVEESIYQ	CCDLAPEARQ	AIRSLIERLY	IGGPLINSKG	QNCGYRRCRA	2700
SGVLTTSCGN	TLICYLKATA	ACRAAKLQDC	TMLVNGDDLV	VICESAGIQE	2750
DAAALRAFTE	AMTRYSAPPG	DPPQPEYDLE	LITSCSSNVS	VAHDASGKRV	2800
YYLTRDPTTP	LARAAWETAR	HTPINSWLGN	IIMYAPTLWA	RMILMIHFFS	2850
ILLAQEQLEK	ALDCQIYGAC	YSIEPLDLPQ	IIERLHGLSA	FTLHSYSPGE	2900 ·
INRVASCLRK	LGVPPLRTWR	HRARSVRAKL	LSQGGRAATC	GRYLFINWAVR	2950
TKLKLTPIPA	ASQLDLSGWF	VAGYSGGDIY	HSLSRARPRW	FPLCLLLLSV	3000
GVGIYLLPNR					3010

FIG. 14H

#2. Strategy for constructing chimeric clone of HCV (pH77CV-J4) which contains the nonstructural region of strain H77 and the structural region of strain HC-J4



- 1. Fragment A, B, C and D; PCR amplification from pCV-H77C or pCV-J4L6S
 - Fragment A; additional *Cla* I site, artificial *Nde* I site induced by a single mutation (C→T at nt 2765 of H77C) and authentic *Eco*47 III site
 - Fragment B and C; eliminated Nde I site by a single mutation within the primers (C→T at nt 9158 of H77C), and fusion PCR with both fragments
 - Fragment D; artificial *Nde* I site induced by 2 point mutations within the primer (T→A at nt 2762 and C→T at nt 2765 of J4L6S)
- 2. TA cloning of PCR products
- 3. Sequence analysis
- 4. Cloning of Fragment A (Cla I-Eco 47III) and Fragment B/C (Hind III-Afl II) with correct sequence into pCV-H77C
- 5. Complete sequence analysis of new cassette vector [pH77CV], into which the structural regions of different genotypes can be inserted.
- 6. Cloning of Fragment-Age I/Xho I (cut out from pCV-J4L6S) and Fragment D (Xho I-Nde I) with correct sequence into the new cassette vector; 3 piece ligation
- 7. Complete sequence analysis of 1a+1b chimera [pH77CV-J4]
- 8. In vitro transcription (within 24 hours of inoculation)
- 9. Percutaneous intrahepatic transfection into chimpanzee

			CATGAATCAC		50
GGAACTACTG	TCTTCACGCA	GAAAGCGTCT	AGCCATGGCG	TTAGTATGAG	100
TGTCGTCCAG	CCTCCAGGAC	CCCCCTCCC	GGGAGAGCCA	TAGICGICIG	150
CGGAACCGGT	GAGTACACCG	GAATTGCCAG	GACGACCGGG	TCCTTTCTTG	200
GATCAACCCG	CTCAATGCCT	GCACATTTGG	GCGIGCCCCC	GCCACACTCC	250
TAGCCGAGTA	GIGITGGGIC	GCGAAAGGCC	TIGIGGIACT	GCCTGATAGG	300
GIGCTIGGGA	GIGCCCCCCCCC	AGGICICGEA	GACCGTGCAC	CATGAGCACG	350
AATCCTAAAC	CTCAAAGAAA	AACCAAACGT	AACACCAACC	GCCGCCCACA	400
GGACGICAAG	TTCCCGGGGGG	GTGGTCAGAT	CGTTGGTGGA	GITTACCIGT	450
TGCCGCGCAG	GGGCCCCAGG	TIGGGIGIGC	GCGCGACTAG	GAAGGCTTCC	500
GAGCGGTCGC	AACCICGIGG	AAGGCGACAA	CCTATCCCAA	AGGCTCGCCG	550
ACCCGAGGGC	AGGGCCTGGG	CTCAGCCCGG	GIACCCTTGG	CCCCTCTATG	600
GCAATGAGGG	CCIGGGGIGG	GCAGGATGGC	TCCTGTCACC	CCCCCCCCCCC	650
CGGCCTAGIT	GGGGCCCCAC	GCACCCCCCG	CGTAGGTCGC	GEAACTTGGG	700
TAAGGICATC	GATACCCTTA	CATGCGGCTT	CCCCGATCTC	ATGGGGTACA	750
TICCGCICGI	cccccccc	CTAGGGGGGG	CTCCCAGGGC	CTTGGCACAC	800
GCIGICCGGG	TICIGGAGGA	CCCCCTCAAC	TATCCAACAG	GGAACTTGCC	850
CGGTTGCTCT	TICICIAICI	TCCTCTTGGC	TCTGCTGTCC	TGTTTGACCA	900
			TGTCCGGGAT		950
ACGAACGACT	GCTCCAACTC	AAGCATTGIG	TATGAGGCAG	CCCACCTGAT	1000
CATGCATACT	CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	TECCCIGIGI	TCAGGAGGGT	AACAGCTCCC	1050
GTTGCTGGGT	AGCGCTCACT	CCCACGCTCG	CCCCACCAA	TECCAGCETC	1100
CCCACTACGA	CAATACGACG	CCACGICGAC	TICCICCITG	GCACGCCTGC	1150
TITCICCICC	GCTATGTACG	TGGGGGATCT	CIGCGGATCT	ATTITICCICG	1200
TCTCCCAGCT	GITCACCTIC	TCGCCTCGCC	CCCATCACAC	AGTGCAGGAC	1250
TGCAACTGCT	CAATCTATCC	CGGCCATGIA	TCAGGTCACC	CATCCCTTG	1300
GGATATGATG	ATGAACTGGT	CACCTACAAC	ACCCTAGIG	GIGICCCAGT	1350
TGCTCCGGAT	CCCACAAGCT	GICGICGACA	TEGTESCESS	GGCCCACTGG	1400
GGAGICCIGG	CCCCCLLCC	CIACIATICC	ATGGTAGGGA	ACTGGGCTAA	1450
GGTTCTGATT	GIGGCGCTAC	TCTTTGCCGG	CGITGACGGG	GAGACCCACA	1500
			CCCGGGTTCAC		1550
			GIGAATACCA		1600
			TGACTCCCTC		1650
			TCAACTCGTC		1700
			TEGITESCCC		1750
CCCCATCACC	TATACTAAGC	CTAACAGCTC	GGATCAGAGG	CCTTATIGCT	1800

FIG. 16A

GGCATTACGC	GCCTCGACCG	TGTGGTGTCG	TACCCGCGTC	GCAGGIGIGI	1850
GGICCAGIGI	ATTGTTTCAC	CCCAAGCCCT	GITGIGGIGG	GGACCACCGA	1900
TCGTTCCCGT	GTCCCTACGT	ATAGCTGGGG	GGAGAATGAG	ACAGACGIGA	1950
TGCTCCTCAA	CAACACGCGT	CCGCCACAAG	GCAACTGGTT	CGGCTGTACA	2000
TCCATCAATA	GIACIGGGIT	CACTAAGACG	TGCGGAGGTC	CCCCGIGIAA	2050
CATCGGGGGG	GICGGIAACC	GCACCTTGAT	CTGCCCCACG	GACTGCTTCC	2100
GGAAGCACCC	CGAGGCTACT	TACACAAAAT	GIGGCICGGG	GCCCTGGTTG	2150
ACACCTAGGT	CCCTAGTAGA	CTACCCATAC	AGGCTTTGGC	ACTACCCCTG	2200
CACTCTCAAT	TTTTCCATCT	TTAAGGTTAG	GAIGIAIGIG	GGGGGGIGG	2250
AGCACAGGCT	CAATGCCGCA	TGCAATTGGA	CTCGAGGAGA	GCGCTGTAAC	2300
TIGGAGGACA	CCCATACCTC	AGAACTCAGC	CCCCTCCTCC	TGICTACAAC	2350
AGAGTGGCAG	ATACTGCCCT	GTGCTTTCAC	CACCCTACCG	GCTTTATCCA	2400
CIGGITIGAT	CCATCTCCAT	CAGAACATOG	TGGACGTGCA	ATACCIGIAC	2450
GGIGIAGGGT	CAGCGITIGI	CICCITICCA	ATCAAATGGG	AGIACATCCT	2500
GITGCITTIC	CTTCTCCTCG	CAGACGCGCG	CGIGIGIGCC	TGCTTGTGGA	2550
TGATGCTGCT	GATACCCCAG	GCTGAGGCCG	CCTTAGAGAA	CITEGIGGIC	2600
CTCAATGCGG	CCICCCIGCC	CGGAGCGCAT	GGIATICICT	CCTTTCTTGT	2650
GITCITCICC	GCCGCCTGGT	ACATTAAGGG	CAGGCTGGCT	CCTGGGGGGG	2700
CGTATCCTTT	TTATGGCGTA	TESCCECTEC	TCCTGCTCCT	ACTGGCGTTA	2750
CCACCACGAG	CATATGCACT	GGACACGGAG	GIGGCCGCGT	CCTCTCCCCC	2800
CGLIGLICLL	GICGGGITAA	TEGECETGAC	TCTGTCGCCA	TATTACAAGC	2850
GCTATATCAG	CIGGIGCATG	TEGIGECTIC	AGIATITICT	GACCAGAGIA	2900
GAAGCGCAAC	TGCACGIGIG	GGTTCCCCCCC	CTCAACGTCC	GGGGGGGGG	2950
CGATGCCGTC	ATCTTACTCA	TGIGIGIAGT	ACACCCGACC	CTGGTATTTG	3000
ACATCACCAA	ACTACTCCTG	GCCATCTTCG	GACCCCTTTG	GATTCTTCAA	3050
GCCAGTTTGC	TTAAAGICCC	CIACTICGIG	CGCGTTCAAG	CCCTTCTCCCG	3100
GATCTGCGCG	CTAGCGCGGA	AGATAGCCGG	AGGICATTAC	GIGCAAAIGG	3150
CCATCATCAA	GTTAGGGGGG	CITACIGGCA	CCIAIGIGIA	TAACCATCIC	3200
ACCCCICTIC	GAGACTGGGC	GCACAACGGC	CIGOGAGAIC	TGGCCGTGGC	3250
			GACCAAGCTC		3300
				CCCCGICICI	3350
			CCACCCGACG		3400
			GGCGTACGCC		3450
GAGGCCTCCT	AGGGIGIATA	ATCACCAGCC	TGACTGGCCG	GCACAAAAAC	3500
				AAACCTTCCT	3550
GGCAACGTGC	ATCAATGGGG	TATGCTGGAC	TGTCTACCAC	GGGGCCGGAA	3600

FIG. 16B

CGAGGACCAT CGCATCACCC AAGGGTCCTG TCATCCAGAT GTATACCAAT	3650
GIGGACCAAG ACCTIGIGGG CIGGCCCGCT CCICAAGGIT CCCGCTCATT	3700
GACACCCIGT ACCIGCGCT CCICGGACCT TIACCIGGIC ACGACGCACG	3750
CCGATGTCAT TCCCGTGCGC CGGCGAGGTG ATAGCAGGGG TAGCCTGCTT	3800
TCGCCCCGC CCATTICCIA CITGAAAGGC TCCTCGGGGG GICCGCTGIT	3850
GIGCCCCCC GGACACCCC TGCGCCIATT CAGGGCCGCG GIGIGCACCC	3900
GIGGAGIGGC TAAAGCGGIG GACITIAICC CIGIGGAGAA CCIAGGGACA	3950
ACCATGAGAT CCCCGGTGTT CACGGACAAC TCCTCTCCAC CAGCAGTGCC	4000
CCAGAGCITC CAGGIGGCCC ACCIGCAIGC TCCCACCGGC AGCGGIAAGA	4050
GCACCAAGGT CCCGGCTGCG TACGCAGCCC AGGGCTACAA GGTGTTGGTG	4100
CTCAACCCCT CIGITGCTGC AACGCTGGGC TTTGGTGCTT ACATGTCCAA	4150
GCCCATGGG GITGATCCTA ATATCAGGAC CGGGGTGAGA ACAATTACCA	4200
CIGGCAGCCC CATCACGIAC TCCACCIACG GCAAGITCCT TGCCGACGGC	4250
COGTOCTCAG GACGICCTTA TCACATAATA ATTIGICACG AGICCCACTC	4300
CACGGATGCC ACATCCATCT TGGGCATCGG CACTGTCCTT GACCAAGCAG	4350
AGACTGCGGG GCCGAGACTG GTTGTGCTCG CCACTGCTAC CCCTCCGGGC	4400
TOOGTCACTG TGTCCCATCC TAACATCGAG GAGGITGCTC TGTCCACCAC	4450
CGGAGAGATC CCCTTTTACG GCAAGGCTAT CCCCCTCGAG GTGATCAAGG	4500
GGGGAAGACA TCTCATCTTC TGCCACTCAA AGAAGAAGTG CGACGAGCTC	4550
GCCGCGAAGC TGGTCGCATT GGGCATCAAT GCCGTGGCCT ACTACCGCGG	4600
TCTTGACGTG TCTGTCATCC CGACCAGCGG CGATGTTGTC GTCGTGTCGA	4650
CCGATGCTCT CATGACTGGC TTTACCGGGG ACTTCGACTC TGTGATAGAC	4700
TGCAACACGT GTGTCACTCA GACAGTCGAT TTCAGCCTTG ACCCTACCTT	4750
TACCATTGAG ACAACCACGC TCCCCCAGGA TGCTGTCTCC AGGACTCAAC	4800
GCCGGGGCAG GACTGGCAGG GCGAAGCCAG GCATCTATAG ATTTGTGGCA	4850
CCGGGGGAGC GCCCTCCGG CATGITCGAC TCGTCCGTCC TCTGTGAGTG	4900
CTATGACGCG GGCTGTGCTT GGTATGAGCT CACGCCCGCC GAGACTACAG	4950
TTAGGCTACG AGCGTACATG AACACCCCGG GGCTTCCCGT GTGCCAGGAC	5000
CATCITGAAT TITIGGGAGGG CGICTTIACG GGCCICACIC ATATAGATGC	5050
CCACTITITIA TCCCAGACAA AGCAGAGIGG GGAGAACTIT CCTTACCIGG	5100
TAGOGTACCA AGCCACOGIG TGOGCTAGGG CTCAAGCCCC TCCCCCATCG	5150
TECCACCAGA TETECAACIG TITCATCOCC CITAAACCCA CCCTCCATCG	5200
GCCAACACCC CTGCTATACA GACTGGGGGC TGTTCAGAAT GAAGTCACCC	5250
TGACGCACCC AATCACCAAA TACATCATGA CATGCATGIC GGCCGACCIG	5300
GAGGICGICA CGAGCACCIG GGIGCICGIT GGCGGCGICC TGCCIGCICT	5350
OCCOCCITAT TOCCTOTICAA CABOCTOCCT OCTCATACTG OCCAGGATCG	5400

FIG. 16C

TCTTGTCCCG	GAAGCCGGCA	ATTATACCIG	ACAGGGAGGT	TCTCTACCAG	5450
CACTICCATG	AGATGGAAGA	GIGCICICAG	CACTTACCGT	ACATOGAGCA	5500
AGGGATGATG	CICGCIGAGC	AGITCAAGCA	GAAGGCCCTC	GCCTCCTGC	5550
AGACCGCGTC	CCGCCATGCA	GAGGITATCA	CCCCTCCTGT	CCAGACCAAC	5600
TGGCAGAAAC	TCGAGGICIT	TTGGGCGAAG	CACATGTGGA	ATTICATCAG	5650
TGGGATACAA	TACTIGGGG	GCCTGTCAAC	GCIGCCIGGT	AACCCCGCCA	5700
TIGCTICATT	GATGGCTTTT	ACAGCIGCCG	TCACCAGCCC	ACTAACCACT	5750
GGCCAAAACCC	TCCTCTTCAA	CATATIGGGG	GGGIGGGIGG	CIGCCCAGCT	5800
acceccaca	GGIGCCGCIA	CIGCCITIGI	GGGIGCIGGC	CTAGCTGGCG	5850
CCGCCATCGG	CAGCGITGGA	CIGGGGAAGG	TCCTCGTGGA	CATTCTTGCA	5900
GGGTATGGCG	CCCCCTCCC	GGGAGCTCTT	GIAGCATICA	AGATCATGAG	5950
CGGIGAGGIC	CCCTCCACGG	AGGACCIGGI	CAATCIGCIG	CCCGCCATCC	6000
TCICGCCIGG	AGCCCTTGTA	GICGGIGIGG	TCTGCGCAGC	AATACTGCGC	6050
CGGCACGITG	GCCCGGGGGGA	GGGGGCAGTG	CAATGGATGA	ACCGGCTAAT	6100
AGCCTTCGCC	TCCCGGGGGA	ACCATGITTC	CCCCACGCAC	TACGTGCCGG	6150
AGAGCGATGC	AGCCGCCCCC	GICACTOCCA	TACTCAGCAG	CCTCACTGIA	6200
ACCCAGCTCC	TGAGGCGACT	GCATCAGTGG	ATAAGCTCGG	AGIGIACCAC	6250
TCCATGCTCC	GGTTCCTGGC	TAAGGGACAT	CTGGGACTGG	ATATGCGAGG	6300
TGCTGAGCGA	CTTTAAGACC	TGGCTGAAAG	CCAAGCTCAT	GCCACAACTG	6350
CCTGGGATTC	CCTTIGIGIC	CTGCCAGCGC	GGGTATAGGG	GGGICIGGCG	6400
AGGAGACGGC	ATTATGCACA	CTCGCTGCCA	CIGIGGAGCT	GAGATCACTG	6450
GACATGICAA	AAACGGGACG	ATGAGGATCG	TCGGTCCTAG	GACCTGCAGG	6500
AACATGTGGA	GIGGGACGIT	CCCCATTAAC	GCCTACACCA	CCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCCC	6550
TACTCCCCTT	CCTGCGCCGA	ACTATAAGTT	CCCCTGTCC	AGGIGICIG	6600
CAGAGGAATA	CGTGGAGATA	AGGCGGGTGG	GGGACTTCCA	CIACGIATCG	6650
GGIATGACTA	CIGACAAICT	TAAATGCCCG	TGCCAGATCC	CATCGCCCGA	6700
ATTTTTCACA	GAATTGGACG	GGGTGCGCCT	ACACAGGITT	CCCCCCTT	6750
GCAAGCCCTT	GCTGCGGGAG	GAGGIATCAT	TCAGAGTAGG	ACTCCACGAG	6800
TACCCCGGTGG	GGTCGCAATT	ACCTTGCGAG	CCCGAACCGG	ACGTAGCCGT	6850
GITGACGICC	ATGCTCACTG	ATCCCTCCCA	TATAACAGCA	CACCCCCCCC	6900
CGAGAACGIT	GGCGAGAGGG	TCACCCCCTT	CTATGGCCAG	CTCCTCGGCT	6950
AGCCAGCIGI	CCCCTCCATC	TCTCAAGGCA	ACTTGCACCG	CCAACCATGA	7000
CTCCCCTGAC	GCCGAGCTCA	TAGAGGCTAA	CCTCCTGTGG	AGGCAGGAGA	7050
TGGGCGGCAA	CATCACCAGG	GTTGAGTCAG	AGAACAAAGT	OGIGATICIG	7100
GACTCCTTCG	ATCCGCTTGT	GGCAGAGGAG	GATGAGCGGG	AGGICICCGT	7150
ACCTGCAGAA	ATTCTGCGGA	AGTCTCGGAG	ATTCGCCCGG	CCCTCCCCC	7200

FIG. 16D

pH77CV-J4 Sequence

```
TCTGGGGGG GCCGGACTAC AACCCCCGC TAGTAGAGAC GTGGAAAAAG
                                                          7250
CCTGACTACG AACCACCTGT GGTCCATGGC TGCCCGCTAC CACCTCCACG
                                                          7300
GICCCCICCI GIGCCICCGC CICGGAAAAA GCGIACGGIG GICCICACCG
                                                          7350
AATCAACCCT ATCTACTGCC TTGGCCGAGC TTGCCACCAA AAGTTTTGGC
                                                          7400
ACCICCICAA CITCOGCAT TACGGGGGAC AATACGACAA CATCCICTGA
                                                          7450
COCCCCCC TCTCCCTCCC CCCCCACTC CCACCTTCAC TCCTATTCTT
                                                          7500
CCATGOCCC CCTGCAGGG CAGOCTGGG ATCCGGATCT CAGCGACGG
                                                          7550
TCATGGTCGA CGGTCAGTAG TGGGGCCCCAC ACGGAAGATG TCGTGTGCTG
                                                          7600
CICAATGICT TATTCCIGGA CAGGOGCACT OGTCACCOCG TGOGCIGOGG
                                                          7650
AAGAACAAAA ACTGCCCATC AACGCACTGA GCAACTCGTT GCTACGCCAT
                                                          7700
CACAATCTGG TGTATTCCAC CACTTCACGC AGTGCTTGCC AAAGGCAGAA
                                                          7750
GAAAGICACA TITGACAGAC TOCAAGITCT GGACAGCCAT TACCAGGACG
                                                          7800
TECTCAAGGA GGTCAAAGCA GCGCCGTCAA AAGTGAAGGC TAACTTGCTA
                                                          7850
TOCGTAGAGG AAGCTTGCAG OCTGACGCCC CCACATTCAG CCAAATCCAA
                                                          7900
GITTGGCTAT GGGGCAAAAG ACGTCCGTTG CCATGCCAGA AAGGCCGTAG
                                                          7950
CCCACATCAA CTCCGTGTGG AAAGACCTTC TGGAAGACAG TGTAACACCA
                                                          8000
ATAGACACTA CCATCATGGC CAAGAACGAG GTTTTCTGCG TTCAGCCTGA
                                                          8050
GAAGGGGGT CGTAAGCCAG CTCGTCTCAT CGTGTTCCCC GACCTGGGCG
                                                          8100
TECECETETE CEAGAACATE COCCIGIACE ACGIGGITAG CAACCICCCC
                                                          8150
CTGGCCGTGA TGGGAAGCTC CTACGGATTC CAATACTCAC CAGGACAGCG
                                                          8200
CGTTGAATTC CTCGTCCAAG CGTCGAAGTC CAAGAAGACC CCGATGCGT
                                                          8250
TOTOGIATGA TACCOGCIGI TITGACICCA CAGTCACTGA GAGCGACATC
                                                          8300
CGIACGGAGG AGCCAATTIA CCAATGITGT GACCTGGACC CCCAAGCCCG
                                                          8350
CGTGCCCATC AAGTCCCTCA CTGAGAGGCT TTATGTTGGG GGCCCTCTTA
                                                          8400
CCAATTCAAG GGGGAAAAC TGGGGCTACC GCAGGTGCCG CGCGAGCGGC
                                                          8450
GTACTGACAA CTAGCTGTGG TAACACCCTC ACTTGCTACA TCAAGGCCCG
                                                          8500
GCCACCTGT CGAGCCCCAG GCTCCAGGA CTGCACCATG CTCGTGTGTG
                                                          8550
GCCACCACTT AGICGITATC TGICAAAGIG CGGGGGTCCA GCACGACGCG
                                                          8600
                                                          8650
COGAGODICA GAGODITCAD GGAGGDATG ACCAGGIACT COGOCOCCC
CGGGGACCCC CCACAACCAG AATACGACTT GGAGCTTATA ACATCATGCT
                                                          8700
CCTCCAACGT GTCAGTCGCC CACGACGCG CTGGAAACAG GGTCTACTAC
                                                          8750
CTTACCCGTG ACCCTACAAC CCCCCTCGCG AGAGCCGCGT GGGAGACAGC
                                                          8800
AAGACACACT CCAGTCAATT CCTGGCTAGG CAACATAATC ATGTTTGCCC
                                                          8850
                                                          8900
CCACACTGTG GGCGAGGATG ATACTGATGA CCCATTTCTT TAGGGTCCTC
                                                          8950
ATACCACC ATCACCTICA ACACCTCTT AACTGTGAGA TCTACCGACC
                                                          9000
CTGCTACTCC ATAGAACCAC TGGATCTACC TCCAATCATT CAAAGACTCC
```

FIG. 16E

pH77CV-J4 Sequence

ATGGCCTCAG	CGCATTTTCA	CTCCACAGIT	ACTCTCCAGG	TGAAATCAAT	9050
AGGGTGGCCG	CATGCCTCAG	AAAACTTGGG	GICCCGCCCT	TCCCACCTTC	9100
GAGACACCGG	GCCCGGAGGG	TCCGCCCTAG	GCTTCTGTCC	AGAGGAGGCA	9150
GGGCTGCTAT	ATGTGGCAAG	TACCICTICA	ACTGGGCAGT	AAGAACAAAG	9200
CTCAAACTCA	CTCCAATAGC	GGCCGCIGGC	CCCCTCCACT	TGTCCCGGTTG	9250
GITCACGGCT	GGCTACAGCG	GGGGAGACAT	TTATCACAGC	GIGICICATG	9300
CCCCCCCCCCCC	CIGGITCIGG	TITIGCCTAC	TOCTGCTCGC	TGCAGGGGTA	9350
GGCATCTACC	TCCTCCCCAA	CCCATGAAGG	TTGGGGTAAA	CACTCCGGCC	9400
TCTTAAGCCA	TTTCCTGTTT	TTTTTTTTT	TTTTTTTTT	TITTICTTT	9450
TTTTTTTCTT	TCCTTTCCTT	CTTTTTTTCC	TITCITITIC	CCTTCTTTAA	9500
TEGTESCICC	AICTIAGCCC	TAGICACGGC	TAGCIGIGAA	AGGICCGIGA	9550
GCCGCATGAC	TGCAGAGAGT	GCTGATACTG	GCCTCTCTGC	AGATCATGT	9599

FIG. 16F

H77CV-J4aa Sequence

10	20	30	40	50	
		1234567890			
		DVKFPGGGQI			50
		PEGRAWAQPG			100
		KVIDILICGF			150
		GCSFSIFLLA			200
		MHTPGCVPCV	-		250
		FCSAMYVGDL	-		300
		DMMMWSPIT			350
		VLIVALLFAG			400
SLFSSGASQK	IQLVNINGSW	HINRIALNON	DSLQIGFFAA	LFYAHKFNSS	4 50
GCPERMASCR	PIDWFAQGWG	PITYTKPNSS	DORPYCWHYA	PRPCGVVPAS	500
QVCGPVYCFT	PSPVVVGITD	RSGVPTYSWG	ENEIDVMLLN	NIRPPQGNWF	550
CCIWMNSTGF	TKTCGGPPCN	IGGVGNRTLI	CPIDCFRKHP	EATYTKCGSG	600
PWLTPRCLVD	YPYRLWHYPC	TLNFSIFKVR	MYVOGVEHRL	NAACIWIRGE	650
RONLEDRDRS	ELSPLLLSTT	EWQILPCAFT	TLPALSIGLI	HLHQNIVDVQ	700
YLYGVGSAFV	SFAIKWEYIL	LLFLLLADAR	VCACLWMMLL	IAQAEAALEN	750
LVVLNAASVA	GAHGILSFLV	FFCAAWYIKG	RLAPGAAYAF	YGVWPLLLLL	800
LALPPRAYAL	DTEVAASCGG	VVLVGLMALT	LSPYYKRYIS	WOMWLQYFL	850
TRVEAQLHVW	VPPLNVRGGR	DAVILLMCVV	HPILVFDITK	LLLAIFGPLW	900
ILQASLLKVP	YFVRVQGLLR	ICALARKIAG	CHYVQMAIIK	LGALTGTYVY	950
NHLTPLRDWA	HNGLRDLAVA	VEPVVFSRME	TKLITWGADT	AACGDIINGL	1000
PVSARRGQEI	LLGPADGMVS	KGWRLLAPIT	AYAQQTRGLL	GCIITSLIGR	1050
DKNQVEGEVQ	IVSTATQTFL	ATCINGVCWI	VYHGAGIRTI	ASPKGPVIQM	1100
YTNVDQDLVG	WPAPQGSRSL	TPCTCGSSDL	YLVTRHADVI	PVRRRGDSRG	1150
SLLSPRPISY	LKGSSGGPLL	CPAGHAVGLF	RAAVCTRGVA	KAVDFIPVEN	1200
LGTTMRSPVF	TDNSSPPAVP	QSFQVAHLHA	PIGSGKSTKV	PAAYAAQGYK	1250
VLVLNPSVAA	TLGFGAYMSK	AHGVDPNIRT	GVRTITIGSP	ITYSTYGKFL	1300
ADGGCSGGAY	DIIICDECHS	TDATSILGIG	TVLDQAETAG	ARLVVLATAT	1350
PPGSVIVSHP	NIEEVALSTT	GEIPFYGKAI	PLEVIKGGRH	LIFCHSKKKC	1400
DELAAKLVAL	GINAVAYYRG	LDVSVIPTSG	DVVVVSTDAL	MIGFIGDFDS	1450
VIDONICVIQ	TVDFSLDPTF	TIETTTLPQD	AVSRTQRRGR	TGRGKPGIYR	1500
FVAPGERPSC	MFDSSVLCEC	YDAGCAWYEL	TPAETTVRLR	AYMNTPGLPV	1550
CQDHLEFWEC	VFTGLTHIDA	HFLSQTKQSG	ENFPYLVAYQ	ATVCARAQAP	1600
PPSWDQMWKC	LIRLKPTLHG	PIPLLYRLGA	VONEVILIHE	ITKYIMICMS	1650
ADLEVVISIV	VLVGGVLAAL	AAYCLSTGCV	VIVGRIVLSG	KPAIIPDREV	1700
LYQEFDEMEE	E CSQHLPYIEQ	GMMLAEQFKQ	KALGLLQTAS	RHAEVITPAV	1750
QTWQKLEVE	WAKHMWNFIS	GIQYLAGLSI	LPGNPAIASL	MAFTAAVTSP	1800
LTTGQTLLFN	ILGGWVAAQI	AAPGAATAFV	GAGLAGAAIG	SVGLGKVLVD	1850
- -				ALVVGVVCAA	

FIG. 16G

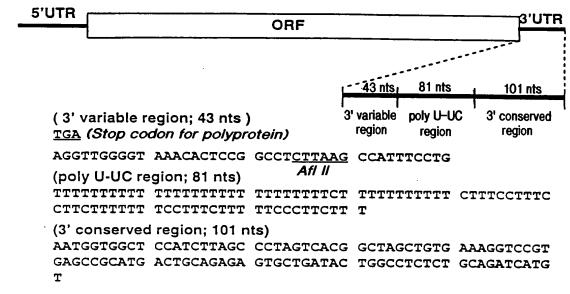
H77CV-J4aa Sequence

					
10	20	30	40	50	
	1234567890				
	GAVQWMNRLI				1950
LIVIQLLRRL	HQWISSECTT	PCSGSWLRDI	WDWICEVLSD	FKIWLKAKIM	2000
PQLPGIPFVS	CQRGYRGVWR	GDGIMHIRCH	CGAETTGHVK	NGIMRIVGPR	2050
TCRNMWSGIF	PINAYTIGPC	TPLPAPNYKF	ALWRVSAEEY	VEIRRVGDFH	2100
YVSGMITIDNL	KCPCQIPSPE	FFTELDGVRL	HRFAPPCKPL	LREEVSFRVG	2150
LHEYPVGSQL	PCEPEPDVAV	LTSMLTDPSH	TTAEAAGRRL	ARGSPPSMAS	2200
SSASQLSAPS	LKATCTANHD	SPDAELIEAN	LLWRQEMGGN	TTRVESENKV	2250
VILDSFDPLV	AEEDEREVSV	PAEILRKSRR	FARALPWAR	PDYNPPLVET	2300
WKKPDYEPPV	VHGCPLPPPR	SPPVPPPRKK	RIVVLTESTL	STALAELATK	2350
SFGSSSTSGI	TGDNITTSSE	PAPSGCPPDS	DVESYSSMPP	LEGEPGDPDL	2400
SDGSWSTVSS	GADTEDVVCC	SMSYSWIGAL	VTPCAAEEQK	LPINALSNSL	2450
LRHHNLVYST	TSRSACQRQK	KVTFDRLQVL	DSHYQDVLKE	VKAAASKVKA	2500
NLLSVEFACS	LTPPHSAKSK	FGYGAKDVRC	HARKAVAHIN	SWKDLLEDS	2550
VTPIDITIMA	KNEVFCVQPE	KGGRKPARLI	VFPDLGVRVC	EKMALYDVV S	2600
KLPLAVMGSS	YGFQYSPGQR	VEFLVQAWKS	KKTPMGFSYD	TRCFDSTVTE	2650
SDIRTEFALY	QCCDLDPQAR	VAIKSLITERL	YVGGPLINSR	GENCGYRRCR	2700
ASGVLTTSCG	NILICYIKAR	AACRAAGLQD	CIMLVCGDDL	VVICESAGVQ	2750
EDAASLRAFT	EAMIRYSAPP	GDPPQPEYDL	ELITSCSSNV	SVAHDGAGKR	2800
VYYLTRDPTT	PLARAAWETA	RHIPVNSWLG	NIIMFAPTLW	ARMILMIHFF	2850
SVLIARDQLE	QALNCEIYGA	CYSIEPLDLP	PIIQRLHGLS	AFSLHSYSPG	2900
EINRVAACLR	KLGVPPLRAW	RHRARSVRAR	LLSRGGRAAI	CGKYLFNWAV	2950
RTKLKLTPIA	AAGRLDLSGW	FTAGYSGGDI	YHSVSHARPR	WFWFCLLLLA	3000
AGVGIYLLPN	R				3011

FIG. 16H

#1a. 3' Deletion mutants of pCV-H77C

Sequence of 3' untranslated region of pCV-H77C



#1a -1. pCV-H77C(-98X); 3' 98 nucleotides removed from pCV-H77C

#1a -2. pCV-H77C(-42X); 3' 42 nucleotides removed from pCV-H77C

#1a -3. pCV-H77C(X-52); All of the 3' UTR sequence, except 3' 49 nucleotides, removed from pCV-H77C

TGAGCCGCAT GACTGCAGAG AGTGCTGATA CTGGCCTCTC TGCAGATCAT
GT

FIG. 17A

#1a -4. pCV-H77C(X); All of the 3' UTR sequence, except 3' 101 nucleotides, removed from pCV-H77C

TGAAATGGTG GCTCCATCTT AGCCCTAGTC ACGGCTAGCT GTGAAAGGTC CGTGAGCCGC ATGACTGCAG AGAGTGCTGA TACTGGCCTC TCTGCAGATC ATGT

#1a -5. pCV-H77C(+49X); The proximal 49 nucleotides of the 3' conserved region (98 nucleotides; AAT not included) removed from pCV-H77C

#1a -6. pCV-H77C(VR-24); First 24 nucleotides of the 3' variable region removed from pCV-H77C

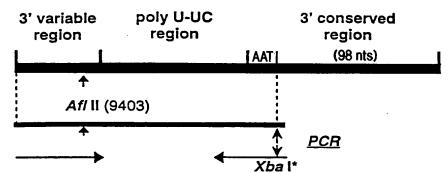
#1a -7. pCV-H77C(-U/UC); Poly U-UC region removed from pCV-H77C

TGAAGGTTGG GGTAAACACT CCGGCCTCTT AAGCCATTTC CTGAATGGTG
GCTCCATCTT AGCCCTAGTC ACGGCTAGCT GTGAAAGGTC CGTGAGCCGC
ATGACTGCAG AGAGTGCTGA TACTGGCCTC TCTGCAGATC ATGT

FIG. 17B

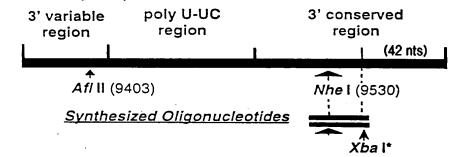
#1b. Strategy of 3' Deletion mutants

#1b -1. pCV-H77C(-98X)

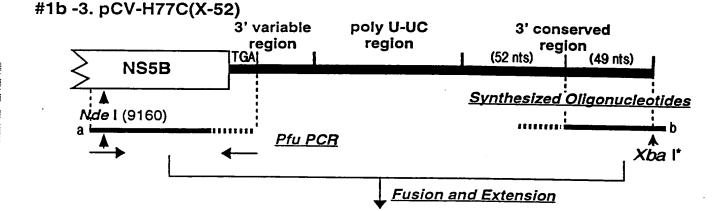


- 1. PCR Amplification
- 2. Purification of PCR products
- 3. Digestion with Afl II and Xba I
- 4. Cloning of Afl II /Xba I fragment into pCV-H77C
- 5. Complete sequence analysis
- 6. in vitro transcription (within 24 hours of inoculation)
- 7. Percutaneous intrahepatic transfection into chimpanzee; 11/26/97 and 12/17/97
- 8. Result : Negative (No replication)

#1b -2. pCV-H77C(-42X)

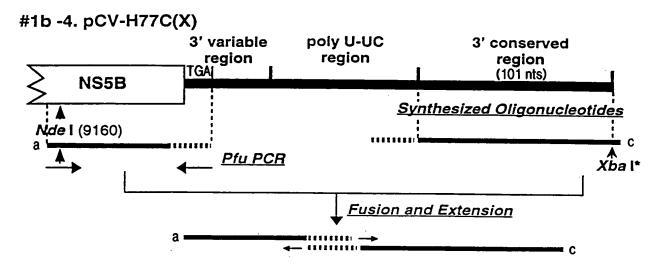


- 1. Synthesis of oligonucleotides (sense and anti-sense)
- 2. Hybridization of oligonucleotides
- 3. Digestion with Nhe I and Xba I
- 4. Cloning of Nhe I /Xba I fragment into pG9-KL26 (3' UTR of H77C)
- 5. Sequence analysis
- 6. Cloning of 3' UTR (-42X) [Afl II /Xba I fragment] into pCV-H77C
- 7. Complete sequence analysis
- 8. in vitro transcription (within 24 hours of inoculation)
- 9. Percutaneous intrahepatic transfection into chimpanzee (Schedule; 1/22/98, 2/5/98)



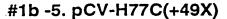
- 1. Fragment a; Pfu PCR amplification and purification
- 2. Fragment b; Synthesized oligonucleotides (anti-sense)
- 3. Fusion and extension
- 4. TA cloning
- 5. Sequence analysis
- 6. Cloning Nde I-Xba I fragment with correct sequence into pCV-H77C
- 7. Complete sequence analysis
- 8. In vitro transcription (within 24 hours of inoculation)
- 9. Percutaneous intrahepatic transfection into chimpanzee

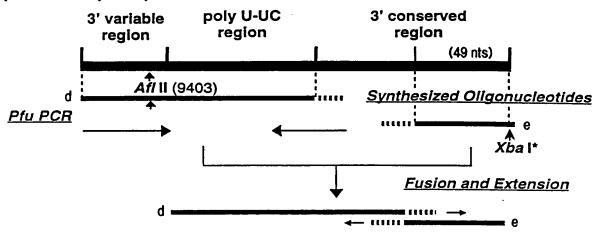
FIG. 17D



- 1. Fragment a; Pfu PCR amplification and purification
- 2. Fragment c; Synthesized oligonucleotides (anti-sense)
- 3. Fusion and extension
- 4. TA cloning
- 5. Sequence analysis
- 6. Cloning Nde I-Xba I fragment with correct sequence into pCV-H77C
- 7. Complete sequence analysis
- 8. In vitro transcription (within 24 hours of inoculation)
- 9. Percutaneous intrahepatic transfection into chimpanzee

FIG. 17E

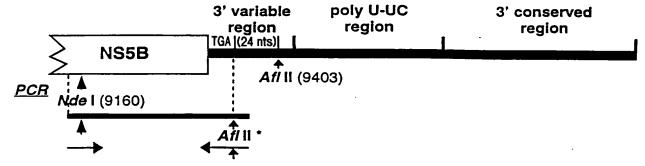




- 1. Fragment d; Pfu PCR amplification and purification
- 2. Fragment e ; Synthesized oligonucleotides (anti-sense)
- 3. Fusion and extension
- 4. TA cloning
- 5. Sequence analysis
- 6. Cloning Afl II-Xba I fragment with correct sequence into pCV-H77C
- 7. Complete sequence analysis
- 8. In vitro transcription (within 24 hours of inoculation)
- 9. Percutaneous intrahepatic transfection into chimpanzee

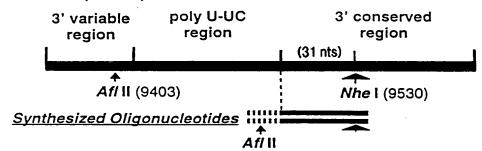
FIG. 17F

#1b -6. pCV-H77C(VR-24)



- 1. PCR Amplification
- 2. Purification of PCR products
- 3. Digestion with Nde I and AfI I
- 4. Cloning of Nde I /Afl II fragment into pCV-H77C
- 5. Complete sequence analysis
- 6. in vitro transcription (within 24 hours of inoculation)
- 7. Percutaneous intrahepatic transfection into chimpanzee

#1b -7. pCV-H77C(-U/UC)



- 1. Synthesis of oligonucleotides (sense and anti-sense)
- 2. Hybridization of oligonucleotides
- 3. Digestion with Aff II and Nhe I
- 4. Cloning of Afl II and Nhe I fragment into pG9-KL26
- 5. Sequence analysis
- 6. Cloning of 3' UTR (-poly U-UC) [Afl II /Xba I fragment] into pCV-H77C
- 7. Complete sequence analysis
- 8. in vitro transcription (within 24 hours of inoculation)
- 9. Percutaneous intrahepatic transfection into chimpanzee

FIG. 17G

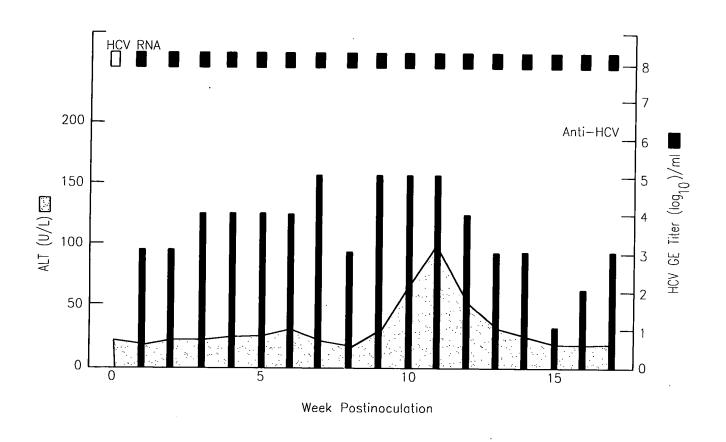


FIG. 18A

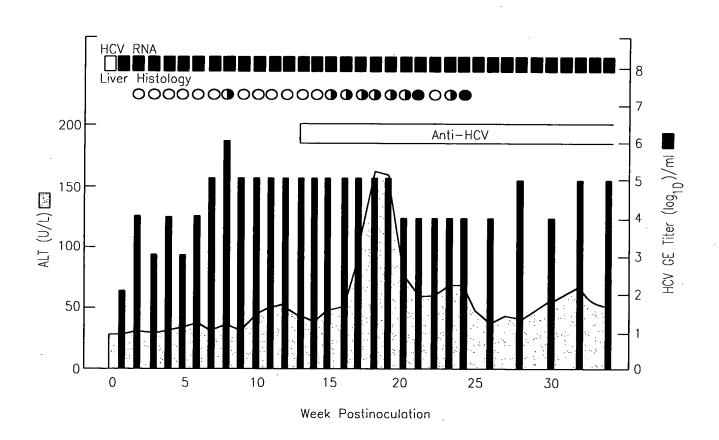


FIG. 18B